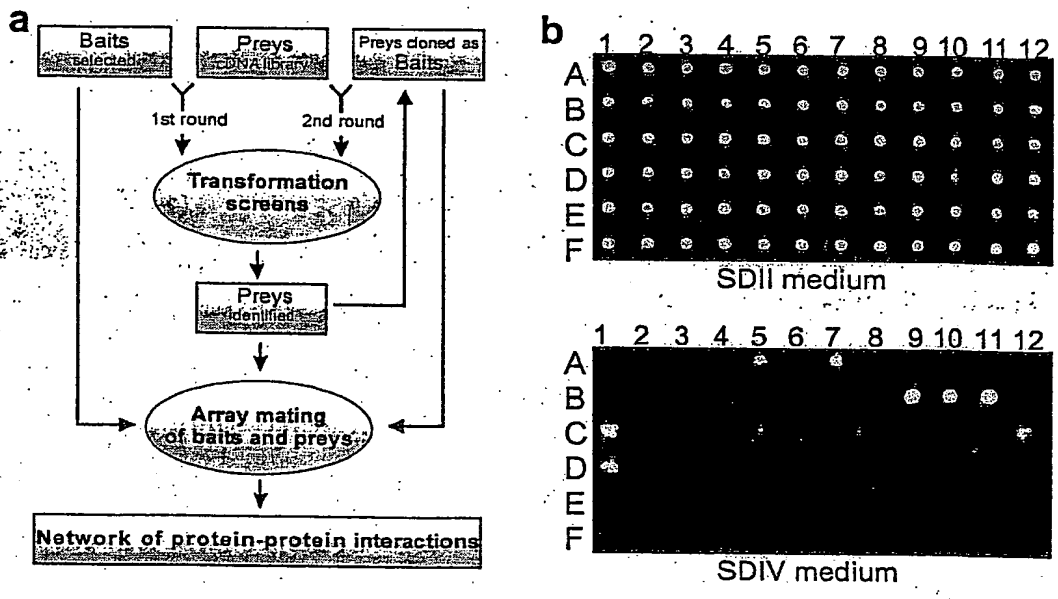
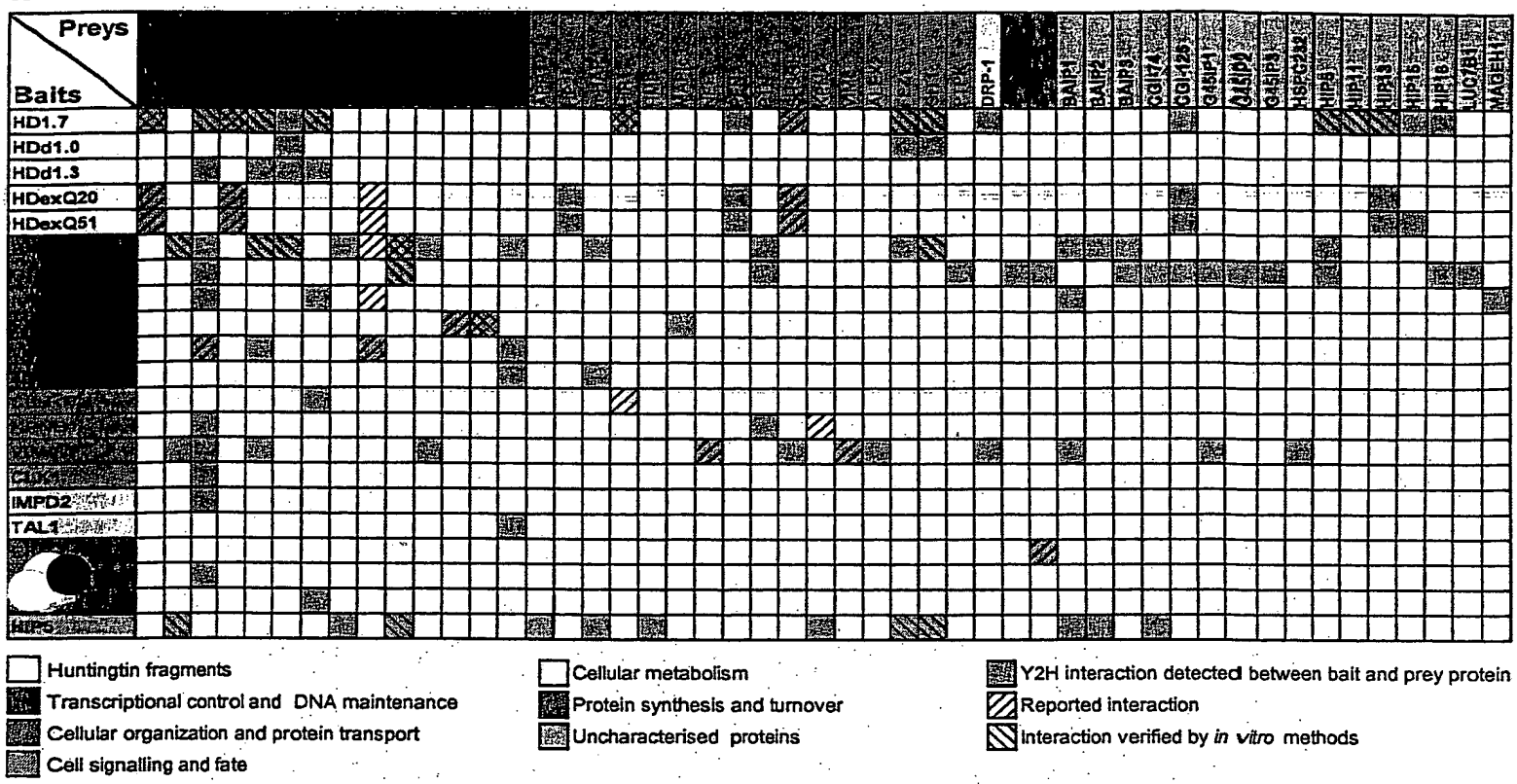


Figure 1

a



b

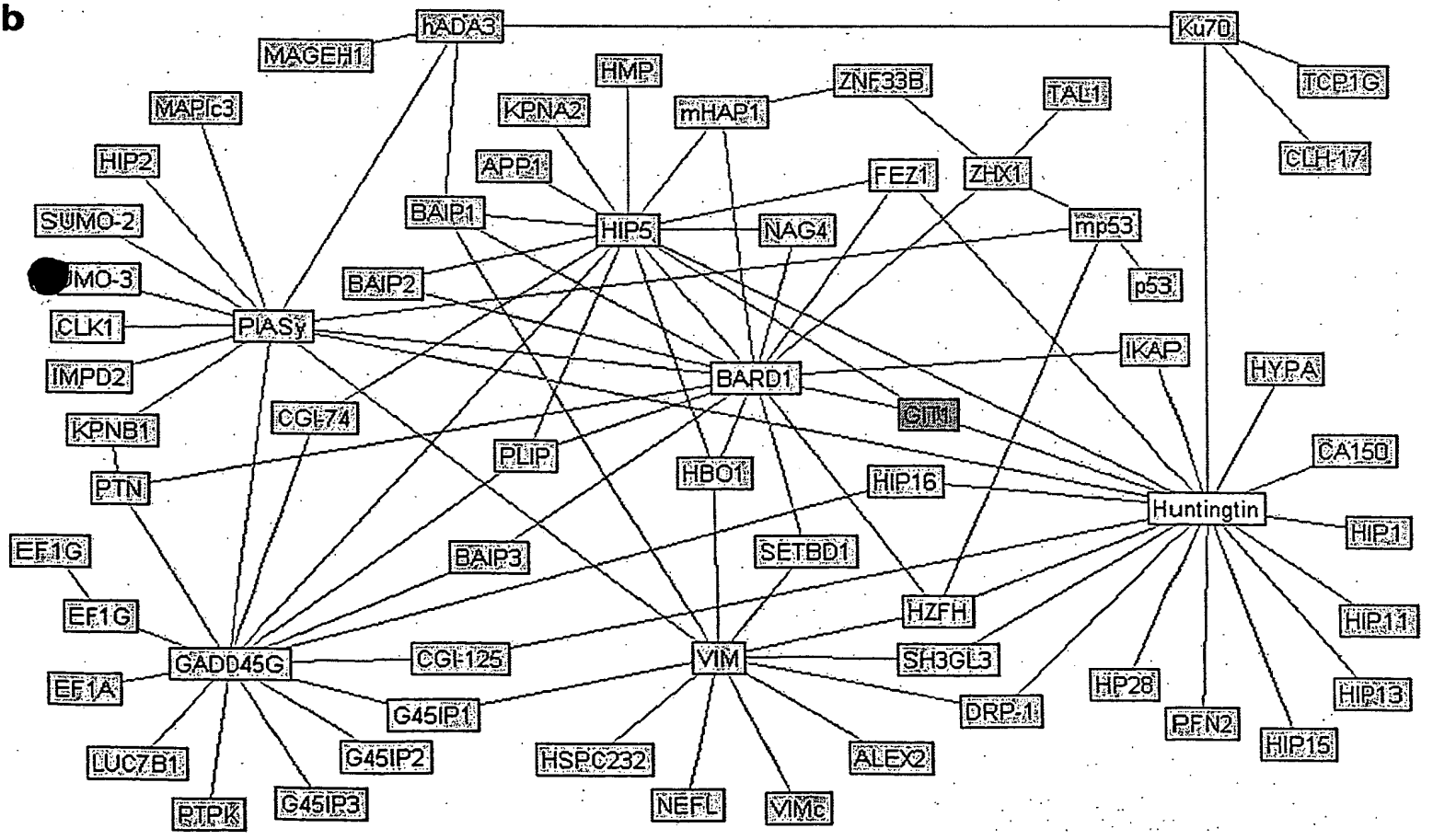


Figure 2

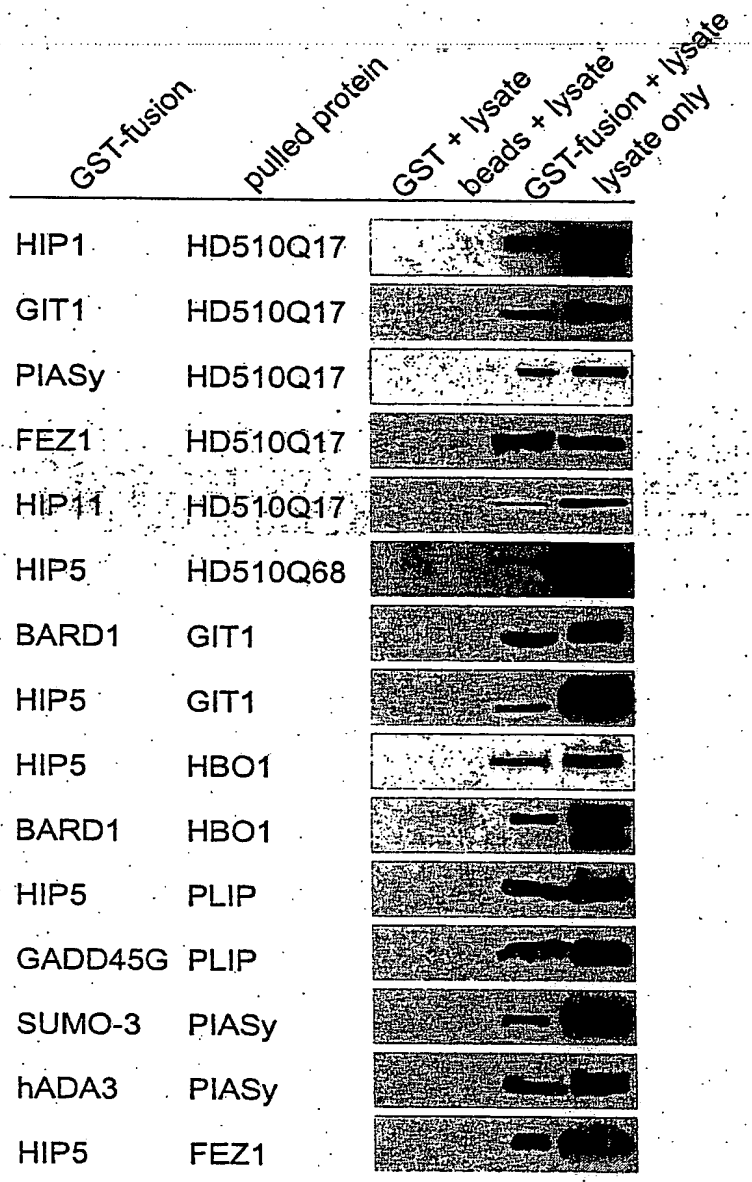


Figure 3

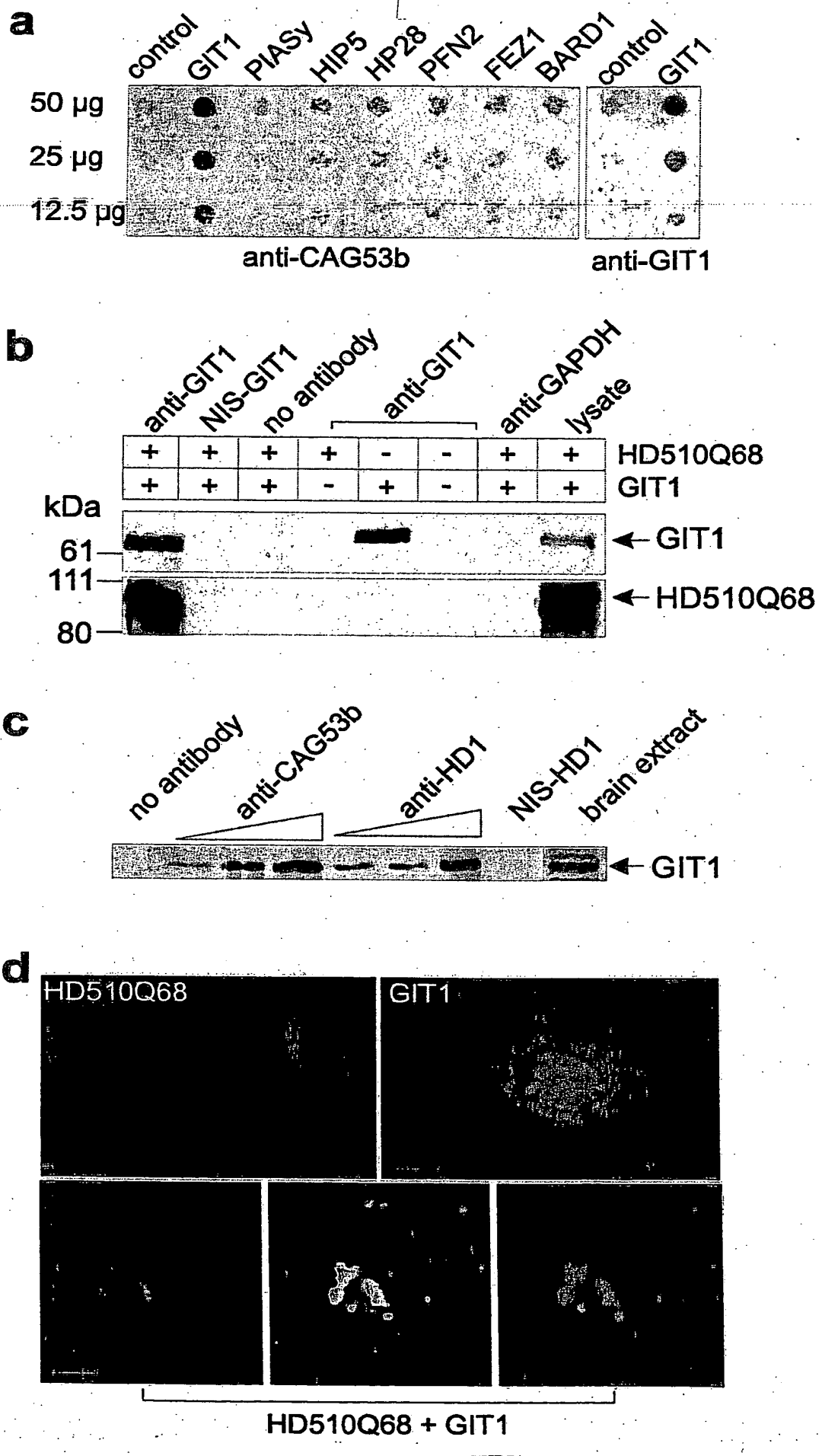


Figure 4

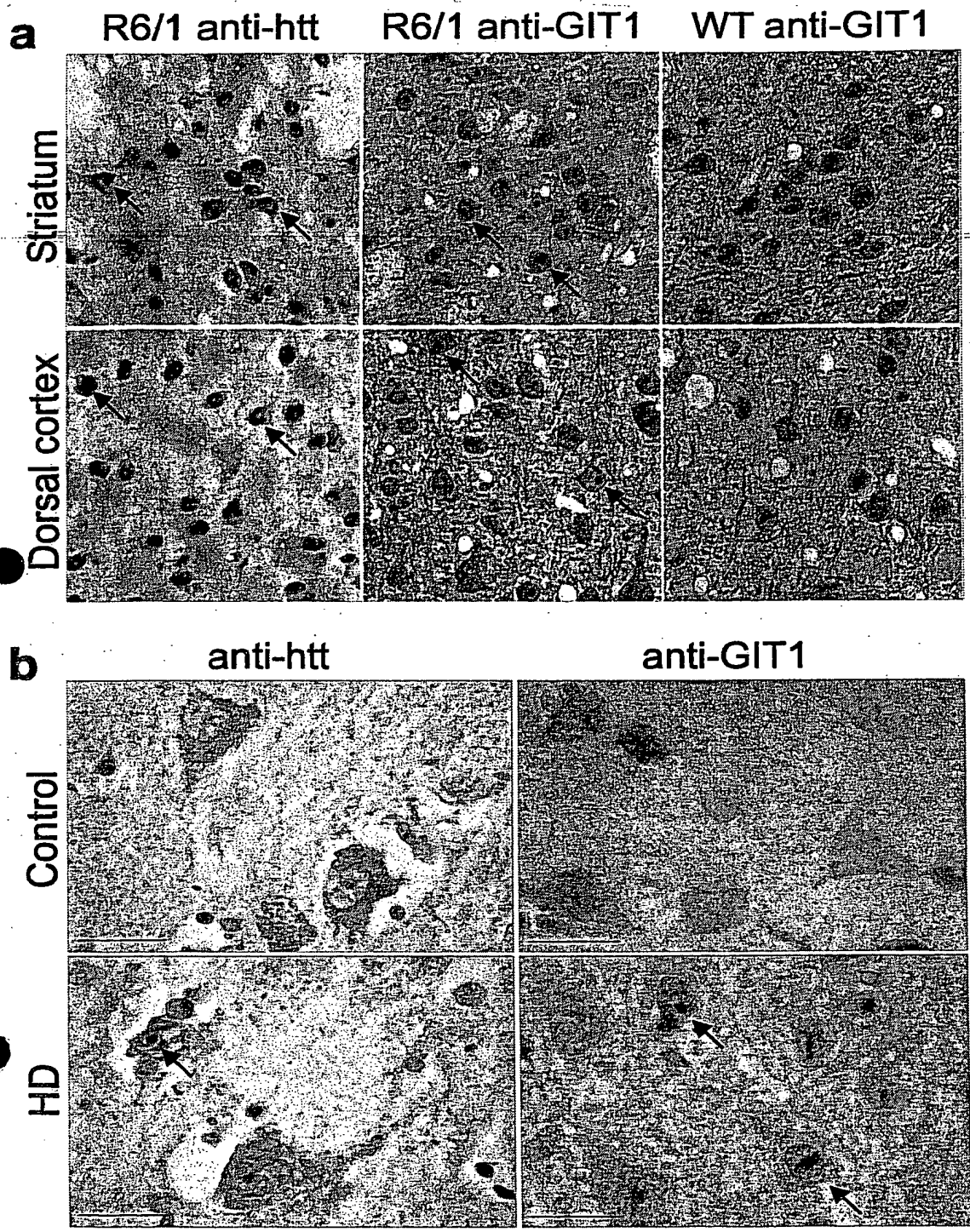


Figure 5

>ALEX2
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VLALLQKSDDPFIQQVALLTLSNNANYSCNQETIRKLGGLPIIANMINKTDPHIKEKALMAMNNLS
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KIKVEILKILSNFAENPDMLKKLLSTQVPASFSSLYNSYVESEILINALTLFEIIYDNLRAEVFNY
REFNKGSLFYLCTTSGVCVKKIRALANHHDLLVKVKVIKLVNKF
>APP1
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THATRVIALINDQRRAALEGFLAALQADPPQAERVLLALRRYLRAEQKEQRHTLRHYQHVAAVDPE
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>BAIP1
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>BAIP2
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>BAIP3
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>CA150
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K

Figure 6

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>CGI-74
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>CLH-17
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EGESQPVKMFDRHSSLAGCQIINYRTDAKQKWLLLTGISAQQNRVVGAMQLYSVDRKVSQPIEGHA
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>CLK1
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EALKHPFFDLLKKSI
>DRP-1
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>EF1G(bait)
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>EF1G(prey)
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Figure 6 (continued)

>G45IP1
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>G45IP2
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RAQPLINLQMVNASLYEHVERMHLIGRRREQLKLLGDYLGLCRSGALKELSKRLNHRNYLLESPHR
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>G45IP3
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NPIYYLRSFFLRRIFLSLLPLQPSPYPPIRRALAPNRHHPAKSPRSPTPKHIRITRIRSINHLSSP
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>GIT1
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IYCQNLCLLAKLFLDHKTLYYDVEPFLFYVMTEADNTGCHLIGYFSKEKNSFLNYNVSCILTMPQY
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Figure 6 (continued)

>HD1.7
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VRPQKCRPYLVNLLPCLTRTSKRPEESVQETLAAAVPKIMASFGNFANDNEIKVLLKAFIANLKSS
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VSSSALTASVKDEISGELAASSGVSTPGSAGHDIITEQPRSQHTAGGLSGSGQL
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>HDd1.3
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>HIP1
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>HIP11
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>HIP13
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>HIP15
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Figure 6 (continued)

>HIP16
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>HIP2
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QYKQNPEMFKQTARLWAHVYAGAPVSSPEYTKKIENLCAMGFDRNAVIVALSSKSWDVETATELLL
SNX
>HIP5 (bait)
FLKSILKKESKYEHGYLKALIINQSFKFGNQKAAAIRDSIELTKEKGAEIPKTIKKLRWFDETSNI
ENNAENSHSLKNKTGTTQQHSQQFHIQSGAGSNIISVSTCAVNSADTKKSREDSISENVTTLGGSG
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TLYCTQRSPVCEESYPSVTLRTAEEESVPLWKRGPNVLHQNKRATGSTVMRRKRIAETKRRNILEQ
KRQNPGSVGQKYSEQINNFGQSVLLSSSEPKQTTRGTSYIEEVSDSTSEFLMAENLVKASVPEDEI
LTVLNSKQIQKSNLPLNKTQQFNICTLSAEEQKILESLNDLNERLHYIQESICKNPSIKNTLQIIP
LLEKREDRTSSCRDKR
>HIP5 (prey)
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ENNAENSHSLKNKTGTTQQHSQQFHIQSGAGSNIISVSTCAVNSADTKKSREDSISENVTTLGGSG
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TLYCTQRSPVCEESYPSVTLRTAEEESVPLWKRGPNVLHQNKRATGSTVMRRKRIAETKRRNILEQ
KRQNPGSVGQKYSEQINNFGQSVLLSSSEPKQTTRGTSYIEEVSDSTSEFLMAENLVKASVPEDEI
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ILTAYASAVGIGTTQVQPE
>HP28
PPADSLLKYDTPVLVSRNTEKRSPKARLLKVSPQQPGPSGSAPQPPKTKLPSTPCVPDPTKQAEEI
LNAILPPREWVEDTQLWIQQVSSTPSTRMDVVHLQEQLDLKLQQRQARETGICPVRRELYSQCFDE
LIREVTINCAERGLLLLRVRDEIRMTIAAYQTLYESSVAFGMRKALQAEQGKSDMERKIAELETEK
RDLERQVNEQKAKCEATEKRESERRQVEEKKHNEEIQFLKRTNQQLKAQLEGIIAPKK
>HSPC232
RRRADGCIYGVSRRARVVAYRRDEMWSEGRYEYERIPRERAPPRSHPSDESGYRWTRDDHSASRQP
EYRDMRDGFRRKSFYSSHYARERSPYKRDNTFFRESPVGRKDSPHSRSGSSVSSRSYSPERSKSYS
FHQSQHRNKERPVQSLKTSRDTSPSSGSAVSSSKVLDKPSRLTEKELAEAASKWAAEKLEKSDESN
LPEISEYEAGSTAPLFTDQPEEPESNTTHGIELFEDSQLTTRSKAIASKTKEIEQVYRQDCETFGM
VVKMLIEKDPSLEKSIQFALRQNLHEIGERCVEELKHFIAEYDTSTQDFGEPF

Figure 6 (continued)

>HYPA
GRRRSSLSPTMRPGTGAERGGLMMGHPGMHYAPMGMHPMGQRANMPPVPHGMMPQMMPPMGGPPMG
QMPGMMSSVMPGMMMSHMSQASMQPALPPGVNSMDVAAGTASGAKSMWTEHKSPDGRTYYYNTETK
QSTWEKPDDLKTPAEQLLSKCPWKEYKSDSGKPYYYNSQTKESRWAKPKELEDLEGYQNTIVAGSL
ITKSNLHAMIKAEESSKQEECTTTSTAPVPTTEIPTTMSTMAAAEAAAAVVAAAAAAAAAAAAANA
NASTSASNTVSGTVPVVPEPEVTSIVATVVDNENTVTLSTEEQAQLTSTPAIQDQSVEVSSNTGEE
TSKQETVADFTPKKEEEESQPAKKTYTWNTKEEAKQAFKELLKEKRVPSNASWEQAMKMIINDPRY
SALAKLSEKKQAFNAYKVQAKKKEKKKKKKK
>HZFH
HARFAEAECLAESHQHLSKESLAGNKPANAVLHKVLNQLEELLSDMKADVTRLPATLSRIPPIAAR
LQMSERSILSRLASKGTEPHPTPAYPPGPYATPPGYGAAFSAAPVGALAAAGANYSQMPAGSFITA
ATNGPPVLVKKEKEMVGALVSDGLDRKEPRAGEVICIDD
>IKAP
LKEGSPLEDLALLEALSEVVQNTENLKDEVYHILKVLFLFEFDEQGRELQKAFEDTLQLMERSLPE
IWTLTYQQNSATPVLGPNSTANSIMASYQQQKTSVPVLDAELFIPPKINRRTQWKLSLLD
>IMPD2
DFLILPGYIDFTADQVDLTSALTKKITLKTPLVSSPMDTVTEAGMAIAMALTGGIGFIHHNCTPEF
QANEVRKVKKYEQGFITDPVVLSPKDRVRDVFEAKARHGFCGIPITDTGRMGSRLVGIISSRDIDF
LKEEEHDCFLEEIMTKREDLVVAPAGITLKEANEILQRSKKGKLPIVNEDDELVAIIARTDLKKNR
DYPLASKDAKKQLLCGAAIGTHEDDKYRLDLLAQAGVDVVVLDSSQGNSIFQINMIKYIKDKYPNL
QVIGGNVVTAAQAKNLIDAGVDALRVGMGSGSICITQEVLACGRPQATAVYKVSEYARRFGVPVIA
DGGIQNVGHIAKALALGASTVMMGSLLAATTEAPGEYFFSDGIRLKKYRGMGSLDAMDKHLSSQNR
YFSEADKIKVAQGVSGAVQDKGSIHKFVPYLIAGIQHSCQDIGAKSLTQVRAMMYSGELKFEKRTS
SAQVEGGVHSLHSYEKRLF
>KPNA2
AWALTNIASGTSEQTKAVVDGGAIPAFISLLASPHAHISEQAVWALGNIAGDGSVFRDLVIKYGAV
DPLLALLAVPDMSSLACGYLRNLTWTLSNLCRNKNPAPPIDAVEQILPTLVRLLHHDDPEVLADTC
WAISYLTDGPNERIGMVVKTGVVPQLVKLLGASELPIVTPALRAIGNIVTGTDEQTQVVIDAGALA
VFPSLLTNPKTNIQKEATWTMSNITAGRQDQIQQVVNHGLVPFLVSVLSKADFKTQKEAVWAVTNY
TSGGTVEQIVYLVHCGIIEPLMNLLTAKDTKIILVILDAISNIFQAAEKLGETEKLSIMIEECGGL
DKIEALQNHENESVYKASLSLIEKYFSVEEEEDQNVVPETTSEGYTFQVQDGAPGTFNF
>KPNB1
LAAVGLVGDLCRALQSNIIPFCDEVMQLLLENLGNENVHRSVKPQILSVFGDIALAIGGEFKKYLE
VVLNTLQQASQAQVDKSDYDMVDYLNELRESCLEAYTGIVQGLKGDQENVHPDVMLVQPRVEFILS
FIDHIAGDEDHTDGVVACAAGLIGDLCTAFGKDVLKLVEARPMIHELLTEGRRSKTNKAKTLATWA
TKELRKLKNQA
>Ku70
KTRTFNTSTGGLLLPSDTKRSQIYGSRQIILEKEETEELKRFDDPGLMLMGFKPLVLLKKHHYLRP
SLFVYPEESLVIGSSTLFSALLIKCLEKEVAALCRYTPRRNIPPYFVALVPQEEELDDQKIQVTPP
GFQLVFLPFADDKRKMPFTEKIMATPEQVGKMKAIVEKLRFTYRSDSFENPVLQQHFRNLEALALD
LMEPEQAVDLTLPKVEAMNKRLGSLVDEFKELVYPPDYNPEGKVTKRKHDNEGSGSKRPKVEYSEE
ELKTHISKGTLGKFTVPMLKEACRAYGLKSGLKKQELLEALTKHFQD
>LUC7B1
VDAVAVDAAAVSAKAEKVHELNEKIGKLLAKAEQLGAEGNVDESQKILMEVEKVRAKKKEAEEEYR
NSMPASSFQQQKLRVCEVCSAYLGLHDNDRRLADHFGGKLHLGFIQIREKLDQLRKTVAEKQEKRN
QDRLRRREEREREERLSRRSGSRTRDRRRSRSRDRRRRRSRSTSRERRKLSRSRSRDRHRRHRSRS
RSHSRGHRRASRDRSAKYKFSRERASREESWESGRSERGPPDWRLESSNGKMASRRSEEKEAGEI

Figure 6 (continued)

>MAGEH1
ASFPRTAVSFEPLAGDMPRGRKSRRRRNARAAEENRNNRKIQASEASETPMAASVVASTPEDDLSG
PEEDPSTPEEASTTPEEASSTAQAQKPSVPRSNFQGTKKSLLMSILALIFIMGNSAKEALVWKVLG
KLGMQPGRQHSIFGDPKKIVTEEFVRRGYLIYKPVPRSSPVEYEFFWGPRAHVESSKLKVMHFVAR
VRNRCSKDWPCNYDWDSDDDAEVEAILNSGARGYSAP
>MAP1lc3
QRRSFADRCKEVQQIRDQHPSKIPVIIERYKGEKQLPVLDKTKFLVPDHVNMSELVKIIRRRLQLN
PTQAFFLLVNQHSMVSVSTPIADIYEQEKDEDGFLYMVYASQETFGF
>mHAP1
PKEQVQSGAGDGTGSGDPAAGTPTTQPAVGPAPEPSAEPKPAPAQGTGSGQKSGSRTKTGSFCRSM
IIGDSDAPWTRYVFQGPYGPRATGLGTGKAEGIWKTPAAYIGRRPGVSGPERAAFIRELQEALCPN
PPPTKKITEDDVKVMLYLLEEKERDLNTAARIGQSLVKQNSVLMEENNKLETMLGSAREEILHLRK
QVNLRDDLLQLYSDSDDDDEEDEEDEEEGEEEEREGQRDQDQQHDHPYGAPKPHPKAETAHRCPQ
LETLQQKLRLLEEENDHLREEASHLDNLEDEEQMLILECVEQFSEASQQMAELSEVLVLRLEGYER
QQKEITQLQAEITKLQQRCQSYGAQTEKLQQMLASEKGIHSESLRAGSYMQDYGSRPRDRQEDGKS
HRQRSSMPAGSVTHYGYSVPLDALPSFPETLAEELRTSLRKFITDPAYFMERRDTHCREGRKKEQR
AMPPPPAX
>mp53
VTETPGPVAPAPATPWPLSSFVPSQKTYQGNYGFHLGFLQSGTAKSVMCTYSPPLNKLFCQLAKTC
PVQLWVSATPPAGSRVRAMAIYKKSQHMTEVVRRCPHHERCSDGDGLAPPQHLIRVEGNLYPEYLE
DRQTFRHSVVVPYEPPEAGSEYTTIHYKYMCNSSCMGGMNRRPILTIITLEDSSGNLLGRDSFEVR
VCACPGRDRRTEEENFRKKEVLCPELPPGSAKRALPTCTSASPPQKKKPLDGEYFTLKIRGRKRFE
MFRELNEALELKDAHATEESGDSRAHSSYLKTKKGQSTSRHKKTMVKKVGPDSD
>NAG4
RDRVENEAEKDLQCHAPVRLDLPPEKPLTSSLAKQEEVEQTPLQEALNQLMRQLQRKDPSAFFSFP
VTDFIAPGYSMIIKHPMDFSTMKEKIKNNDYQSIEELKDNFKLMCTNAMIYNKPETIYYKAAKKLL
HSGMKILSQERIQSLKQSIDFMADLQKTRKQKDGTDTSQSGEDGGCWQRERDSGDAEAHAFKSPS
KENKKKDKDMLEDKFKSNNLEREQEQLDRIVKESGGKLTRRLVNSQCEFERRKPDGTTTLGLLHPV
DPIVGEPGYCPVRLGMTTGRLQSGVNTLQGFKEDKRNKVTPVLYLNYGPYSSYAPHYDSTFANISK
DDSDLIYSTYGEDSDLPSDFSIHEFLATCQDYPYVMADSLLDVLTKGGHSRTLQEMEMSLPEDEGH
TRTLDTAKEMEITEVEPPGRLDSSTQDRLIALKAVTNFGVPVEVFDSEEAEIFQKKLDETTRLLRE
LQEAQNERLSTRPPPNMICLLGPSYREMHLAEQVTNNLKELAQQVTPGDIVSTYGVRKAMGISIPS
PVMENNFVDLTEDTEEPKKTDVAECGPGGS
>NEFL
LSPLSSLSGLPPPPRAGEPPAATMSSFSYEPYYSTSYKRRYVETPRVHISSVRSGYSTARSAYSSY
SAPVSSSLSVRRSYSSSSGSLMPSLENLDLSQVAAISNDLKSIRTQEKAQLQDLNDRFASFIERVH
ELEQQNKVLEAELLVLRQKHSEPSRFRALYEQEIRDLRLAAEDATNEKQALQGEREGLEETLRNLQ
ARYEEEVLSREDAEGRLMEARKGADEAALARAELEKRIDSLMDEISFLKKVHEEEIAELQAQIQYA
QISVEMDVTKPDLSAALKDIRAQYEKLAAKNMQNAEEWFKSRFTVLTESAAKNTDAVRAAKDEVSE
SRRLLKAKTLEIEACRGMNEALEKQLQELEDKQNADISAMQDTINKLENELRTTKSEMARYLKEYQ
DLLNVKMALDIEIAAYRKLLEGEETRLSFTSVGSITSGYSQSSQVFGRSAYGGLQTSSYLMSTRSF
PSYYTSHVQEEQIEVEETIEAAKAEEAKDEPPSEGEAEEEEKDKEEAEEEEAAEEEEAAKEESEEA
KEEEEGGEGEEGEETKEAEEEEKKVEGAGEEQAAKKKD
>p53
MEEPQSDPSVEPPLSQETFSDLWKLLPENNVLSPLPSQAMDDLMLSPDDIEQWFTEDPGPDEAPRM
PEAAPPVAPAPAAPTPAAPAPAPSWPLSSSVPSQKTYQGSYGFRLGFLHSGTAKSVTCTYSPALNK
MFCQLAKTCPVQLWVDSTPPPGTRVRAMAIYKQSQHMTEVVRRCPHHERCSDSDGLAPPQHLIRVE
GNLRVEYLDDRNTFRHSVVVPYEPPEVGSDCTTIHYNYMCNSSCMGGMNRRPILTIITLEDSSGNL
LGRNSFEVRVCACPGRDRRTEEENLRKKGEPHHELPPGSTKRALPNNTSSSPQPKKKPLDGEYFTL
QIRGRERFEMFRELNEALELKDAQAGKEPGGSRAHSSHLKSKKGQSTSRHKKLMFKTEGPDSD

Figure 6 (continued)

>PFN2
APRRPRCSAKGSKMAGWQSYVDNLMCDGCCQEAAIVGYCDAKYVWAATAGGVFQSITPIEIDMIVG
KDREGFFTNGLTLGAKKCSVIRDSLYVDGDCTMDIRTKSQGGEPTYNVAVGRAGRVLVFVMGKEGV
HGGGLNKKAYSMAKYLRDSGF
>PIASy(bait)
LVEAKNMVMSFRVSDLQMLLGFVGRSKSGLKHELVTRALQLVQFDCSPELFKKIKELYETRYAKKN
SEPAPQPHRPLDPLTMHSTYDRAGAVPRTPLAGPNIDYPVLYGKYLNGLGRLPAKTLKPEVRLVKL
PFFNMLDELLKPTELVPQNNEKLQESPCIFALTPRQVELIRNSRELQPGVKAVQVVLRICYSDTSC
PQEDQYPPNIAVKVNHSYCSVPGYYPSNKPGVEPKRPCRPINLTHLMYLSSATNRITVTWGNYGKS
YSVALYLVRQLTSSELLQRLKTIGVKHPELCKALVKEKLRLDPDSEIATTGVRVSLICPLVKMRLS
VPCRAETCAHLQCFDAVFYLQMNEKKPTWMCPVCDKPAPYDQLIIDGLLSKILSECEDADEIEYLV
DGSWCPIRAEKERSCSPQGAILVLGPSDANGLLPAPSVNGSGALGSTGGGGPVGSMENGKPGADVV
DLTLDSSSSSEDEEEEEEEEEDEDEEGPRPKRRCPFQKGLVPAC
>PIASy(prey)
LVEAKNMVMSFRVSDLQMLLGFVGRSKSGLKHELVTRALQLVQFDCSPELFKKIKELYETRYAKKN
SEPAPQPHRPLDPLTMHSTYDRAGAVPRTPLAGPNIDYPVLYGKYLNGLGRLPAKTLKPEVRLVKL
PFFNMLDELLKPTELVPQNNEKLQESPCIFALTPRQVELIRNSRELQPGVKAVQVVLRICYSDTSC
PQEDQYPPNIAVKVNHSYCSVPGYYPSNKPGVEPKRPCRPINLTHLMYLSSATNRITVTWGNYGKS
YSVALYLVRQLTSSELLQRLKTIGVKHPELCKALVKEKLRLDPDSEIATTGVRVSLICPLVKMRLS
VPCRAETCAHLQCFDAVFYLQMNEKKPTWMCPVCDKPAPYDQLIIDGLLSKILSECEDADEIEYLV
DGSWCPIRAEKERSCSPQGAILVLGPSDANGLLPAPSVNGSGALGSTGGGGPVGSMENGKPGADVV
DLTLDSSSSSEDEEEEEEEEEDEDEEGPRPKRRCPFQKGLVPAC
>PLIP
GEIIEGCRLPVLRRNQDNEDEWPLAEILSVKDISGRKLFYVHYIDFNKRLDEWVTHERLDLKKIQF
PKKEAKTPTKNGLPGSRPGSPEREVKRKVEVVSPATPVPSETAPASVFPQNGAARRAVAAQPGRKR
KSNCLGTDEDSQDSSDGIPSAPRMTGSLVSDRSHDDIVTRMKNIECIELGRHRLKPWYFSPYPQEL
TTLPVLYLCEFCLKYGRSLKCLQRHLTKCDLRHPPGNEIYRKGTISFFEIDGRKNKSYSQNLCLLA
KCFLDHKTLYYDTDPFLFYVMTEYDCKGFHIVGYFSKEKESTEDYNVACILTLPPYQRRGYGKLLI
EFSYELSKVEGKTGTPEKPLSDLGLLSYRSYWSQTILEILMGLKSESGERPQITINEISEITSIKK
EDVISTLQYLNLINYYKGQYILTLSEDIVDGHERAMLKRLLRIDSKCLHFTPKDWSKRGKW
>PTN
LSQRQDQVPRLPVQKSRQESPRAEENPKWREGKKETSESSVQKAGRAAAAQAGAAASRVPGLSGSN
LAPCNKGRLSAREDVSNSKMQAQQYQQQRRKFAAAFLAFIFILAAVDTAEAGKKEKPEKKVKKSDC
GEWQWSVCVPTSGDCGLGTREGTRTGAECKQTMKTQRCKIPCNWKKQFGAECKYQFQAWGECDLNT
ALKTRTGSLKRALHNAECQKTVTISKPCGKLTKPKPQAESKKKKKEGKKQEKMLD
>PTPK
SNYINAALMDSYRQPAAFIVTQYPLPNTVKDFWRLVYDYGCTSIVMLNEVDLSQGCPQYWPEEGML
RYGPIQVECMSCSMDCDVINRIFRICNLTRPQEGYLMVQQFQYLGWASHREVPGSKRSFLKLILQV
EKWQEECEEGEGRTIIHCLNGGGRSGMFCAIGIVVEMVKRQNVVDVFHAVKTLRNSKPNMVEAPEQ
YRFCYDVALEYLESS
>SETBD1
KASTSGLGIKDEGDIKQAKKEDTDDRNKMSVVTESSRNYGYNPSPVKPEGLRRPPSKTSMHQSRRL
MASAQSNPDDVLTLSSSTESEGESGTSRKPTAGQTSATAVDSDDIQTISSGSEGDDFEDKKNMTGP
MKRQVAVKSTRGFALKSTHGIAIKSTNMASVDKGESAPVRKNTRQFYDGEESCYIIDAKLEGNLGR
YLNHSCSPNLFVQNVFVDTHDLRFPWVAFFASKRIRAGTELTWDYNYEVGSVEGKELLCCCGAIEC
RGRLL

Figure 6 (continued)

>SH3GL3
VAGLKKQFHKASQLFSEKISGAEGTKLDDEFLDMERKIDVTNKVVAEILSKTTEYLQPNPAYRAKL
GMLNTVSKIRGQVKTTGYPQTEGLLGDCMLKYGKELGEDSTFGNALIEVGESMKLMAEVKDSLDIN
VKQTFIDPLQLLQDKDLKEIGHHLKKLEGRRLDYDYKKKRVGKIPDEEVRQAVEKFEESKELAERS
MFNFLENDVEQVSQLAVFIEAALDYHRQSTEILQELQSKLQMRISAASSVPRREYKPRPVKRSSSE
LNGVSTTSVVKTTGSNIPMDQPCCRGLYDFEPENQGELGFKEGDIITLTNQIDENWYEGMIHGESG
FFPINYVEVIVPLPQ
>SUMO-2
RPRAQLRRESGGAESVTRPLRAASPAPPPRAARAAMSEEKPKEGVKTENDHINLKVAGQDGSVVQF
KIKRHTPLSKLMKAYCERQGLSMRQIRFRFDGQPINETDTPAQLEMEDEDTIDVFQQQTGGVPESS
LAGHSF
>SUMO-3
PSSTAAASFFCRSWCCLCARLVRTWYLFCEAAAEETPALAMADEKPKEGVKTENNDHINLKVAGQD
GSVVQFKIKRHTPLSKLMKAYCERQGLSMRQIRFRFDGQPINETDTPAQLEMEDEDTIDVFQQQTG
GVY
>TAL1
SSPVKRQRMESALDQLKQFTTVVADTGDFHAIDEYKPQDATTNPSLILAAAQMPAYQELVEEAIAY
GRKLGGSQEDQIKNAIDKLFVLFGAEILKKIPGRVSTEVDARLSFDKDAMVARARRLIELYKEAGI
SKDRILIKLSSTWEGIQAGKELEEQHGIHCNMTLLFSFAQAVACAEAGVTLISPFVGRILDWHVAN
TDKKSYEPLEDPGVKSVTKIYNYYKKFSYKTIVMGASFRNTGEIKALAGCDFLTISPKLLGELLQD
NAKLVPVLSAKAAQASDLEKIHLDEKSFRWLHNEDQMAVEKLSDGIRKFAADAVKLERMLTERMFN
AENGK
>TCPG
QTDIEITREEDFTRILQMEEEYIQQLCEDIIQLKPDVVITEKGISDLAQHYLMRANITAIRRVRKT
DNNRIARACGARIVSRPEELREDDVGTGAGLLEIKKIGDEYFTFITDCKDPKACTILLRGASKEIL
SEVERNLQDAMQVCRNVLLDPQLVPGGGASEMAVAHALTEKSKAMTGVEQWPYRAVAQALEVIPRT
LIQNCGASTIRLLTSLRAKHTQENCETWGVNGETGTLVDMKELGIWEPLAVKLQTYKTAVETAVLL
LRIDDIVSGHKKKGDDQSRQGGAPDAGQE
>VIM
SPRQRRSRAPTTHTHRALVRLFSGSQSAPPPPPRPSPPSAAMSTRSVSSSSYRRMFGGPGTASRPS
SSRSYVTTSTRTYSLGSALRPSTSRSLYASSPGGVYATRSSAVRLRSSVPGVRLLQDSVDFSLADA
INTEFKNTRTNEKVELQELNDRFANYIDKVRFLEQQNKILLAELEQLKGQGKSRLGDLYEEEMREL
RRQVDQLTNDKARVEVERDNLAEDIMRLREKLQEEMLQREEAENTLQSFRQDVDNASLARLDLERK
VESLQEEIAFLKKLHEEEIQELQAQIQEQHVQIDVDVSKPDLTAALRDVRQQYESVAAKNLQEAEE
WYKSKFADLSEAANRNNDALRQAKQESTEYRRQVQSLTCEVDALKGTNESLERQMREMEENFAVEA
ANYQDTIGRLQDEIQNMKEEMARHLREYQDLLNVKMALDIEIATYRKLLEGEESRISLPLPNFSSL
NLRETNLDSLPLVDTHSKRTLLIKTVETRDGQVINETSQHHDDLE
>VIMc
QEEMLQREEAENTLQSFRQDVDNASLARLDLERKVESLQEEIAFLKKLHEEEIQELQAQIQEQHVQ
IDVDVSKPDLTAALRDVRQQYESVAAKNLQEAEEWYKSKFADLSEAANRNNDALRQAKQESTEYRR
QVQSLTCEVDALKGTNESLERQMREMEENFAVEAANYQDTIGRLQDEIQNMKEEMARHLREYQDLL
NVKMALDIEIATYRKLLEGEESRISLPLPNFSSLNLRETNLDSLPLVDTHSKRTLLIKTVETRDGQ
VINETSQHHDDLE

Figure 6 (continued)

>ZHX1
EQTINDLTFDGSFVKEENAEQAESTEVSSSGISISKTPIMKMMKNKVENKRIAVHHNSVEDVPEEK
ENEIKPDREEIVENPSSSASESNTSTSIVNRIHPSTASTVVTPAAVLPGLAQVITAVSAQQNSNLI
PKVLIPVNSIPTYNAALDNNPLLLNTYNKFPYPTMSEITVLSAQAKYTEEQIKIWFSAQRLKHGVS
WTPEEVEEARRKQFNGTVHTVPQTITVIPTHISTGSNGLPSILQTCQIVGQPGLVLTQVAGTNTLP
VTAPIALTVAGVPSQNNIQKSQVPAAQPTAETKPATAAVPTSQSVKHETALVNPDSFGIRAKKTKE
QLAELKVSYLKNQFPHDSEIIRLMKITGLTKGEIKKWFSDTRYNQRNSKSNQCLHLNNDSSTTIII
DSSDETTESPTVGTAQPKQSWNPFPDFTPQKFKEKTAEQLRVLQASFLNSSVLTDEELNRLRAQTK
LTRREIDAWFTEKKKSKALKEEKMEIDESNAGSSKEEAGETSPADESGAPKSGSTGKICKKTPEQL
HMLKSAFVRTQWPSPEEYDKLAKESGLARTDIVSWFGDTRYAWKNGNLKWYYYYQSANSSSMNGLS
SLRKRGRGRPKGRGRGRPRGRPRGSKRINNWDRGPSLIKFKTGTAILKDYYLKHKFLNEQDLDELV
NKSHMGYEQVREWFAERQRRSELGIELFEENEEEDEVIDDQEEDEEETDDSDTWEPPRHVKRKLSK
SDD
>ZNF33B
CYECGKTFCLKSDLTIHQRTHTGEKPFACPECGKFFSHKSTLSQHYRTHTGEKPYECHECGKIFYN
KSYLTKHNRTHTGEKPYECNECGKTFCQKSQLTQHQRIHIGEKPYECNECGKAFCHKSALIVHQRT
HTQEKPYKCNECGKSFCVKSGLILHERKHTGEKPYECNECGKSFSHKSSLTVHYRAHTGEKSCQCN
ECGKIFYRKSDLAKHQRSHTGEKPYECNTCRKTFSQKSNLIVHQRTHIGEKPYE

Figure 6 (continued)

>ALEX2
GCCGAATCAGTAGTTGGGGCTGCAATGGCTTCTGCAATAGCACCACCTCCCGGGGTGACAGAGGCC
CTTGGGGCTGCAGAAGCCCCTGCAATGGCAGGGGCTCCCAAAGTGGCAGAAGCTCCCAGAGAAGCG
GAGACTTCCAGGGCAGCGGTGCCTCCTGGGACAGTGGTGCCTACCGAAGCGGCAGCACCCACTGAG
GTGACCGAGGGTCCTGGGGTAGCAGCACCTACCAAGGTAGCTGAAGCTCCCGGGGTGGCATCGCCT
ACCGAGGCAGCTGAGGCTCCTGTGCCCGCAACGCCTACTGGGGCTGCAGCACCTACTGGGGCTGCA
GAGTCTCCTGGAACTTCTGGTTCCCCTAGAACAGCGGTGGTTCCTGGAACATCAGCTGCCAAGAAA
GCAACCCCTGGGGCTCACACTGGGGCTATACCGAAAGCCACATCAGCGACTGGAGCGGTACCCAAA
GGTGGAGGCAAGGGTGTAACCAGGTCCCGGAATGGGGGCAAGGGCAAGGGCAAGAAAAGCAAAGTT
GAAGTAGACGAACTGGGGATGGGCTTCCGTCCTGGAGATGGGGCTGCAGCAGCTGCTGCAGCCTCT
GCTAATGGCGGACAGGCTTTCCTGGCAGAGGTCCCTGATTCTGAGGAAGGGGAGTCCGGGTGGACT
GACACAGAGTCAGATTCAGACTCTGAGCCCGAGACCCAGCGCAGAGGGAGGGGAAGAAGACCCGTT
GCCATGCAGAAGCGCCCCTTTCCTTATGAAATTGATGAGATTCTGGGTGTCCGCGATCTCAGGAAG
GTCCTTGCCTTGCTTCAGAAATCTGATGATCCTTTCATCCAACAGGTAGCTTTGCTCACTCTGAGC
AACAATGCCAATTATTCATGCAATCAAGAGACAATCCGCAAATTGGGAGGCCTCCCAATTATTGCA
AACATGATCAACAAAACTGATCCACACATTAAGGAAAAAGCCTTAATGGCCATGAATAACCTGAGT
GAGAATTATGAAAATCAGGGCCGGCTTCAGGTGTACATGAATAAAGTGATGGATGATATCATGGCC
TCTAACCTGAACTCAGCAGTTCAAGTAGTTGGACTAAAATTTCTAACAAACATGACTATTACTAAT
GACTACCAACACCTGCTTGTCAATTCCATTGCAAACTTTTTCCGTTTGCTATCTCAGGGAGGTGGA
AAAATCAAGGTTGAGATTTTGAAAATCCTTTCGAATTTTGCTGAAAATCCAGATATGTTGAAGAAA
CTTCTCAGTACCCAAGTGCCAGCATCATTTAGTTCCCTCTATAATTCTTACGTGGAATCAGAAATC
CTTATTAATGCCCTTACTCTATTTGAGATTATCTATGACAATCTCAGAGCAGAAGTGTTTAACTAT
AGAGAATTCAATAAAGGTTCCCTTTTTTACTTATGCACTACATCTGGAGTGTGTGTTAAGAAAATT
AGAGCCTTAGCAAATCACCATGACCTCTTAGTGAAAGTGAAAGTTATAAAACTAGTGAACAAATTC
>APP1
GAGGAAGAGGAGGAATCCTTCCCACAGCCAGTAGATGATTACTTCGTGGAGCCTCCGCAGGCTGAA
GAGGAAGAGGAAACGGTCCCACCCCCAAGCTCCCATACACTTGCAGTGGTCGGCAAAGTCACTCCC
ACCCCGAGGCCCACAGACGGTGTGGATATTTACTTTGGCATGCCTGGGGAAATCAGTGAGCACGAG
GGGTTCCTGAGGGCCAAGATGGACCTGGAGGAGCGTAGGATGCGCCAGATTAATGAGGTGATGCGT
GAATGGGCCATGGCAGACAACCAGTCCAAGAACCTGCCTAAAGCCGACAGACAGGCCCTGAATGAG
CACTTCCAGTCCATTCTGCAGACTCTGGAGGAGCAGGTGTCTGGTGAGCGACAGCGCCTGGTGGAA
ACCCACGCCACCCGCGTCATCGCCCTTATCAACGACCAGCGCCGGGCTGCCTTGGAGGGCTTCCTG
GCAGCCCTGCAGGCAGATCCGCCTCAGGCGGAGCGTGTCCTGTTGGCCCTGCGGCGCTACCTGCGT
GCGGAGCAGAAGGAACAGAGGCACACGCTGCGCCACTACCAGCATGTGGCCGCCGTGGATCCCGAG
AAGGCACAGCAGATGCGCTTCCAGGTGCATACCCACCTTCAAGTGATTGAGGAGAGGGTGAATCAG
AGCCTGGGCCTGCTTGACCAGAACCCCCACCTGGCTCAGGAGCTGCGGCCCCAAATCCAGGAACTC
CTCCACTCTGAACACCTGGGTCCCAGTGAATTGGAAGCCCCTGCCCCTGGGGGCAGCAGCGAGGAC
AAGGGTGGGCTGCAGCCTCCAGATTCCAAGGATGACACCCCCATGACCCTTCCAAAAGGGTCCACA
GAACAAGATGCTGCATCCCCTGAGAAAGAGAAGATGAACCCGCTGGAACAGTATGAGCGAAAGGTG
AATGCGTCTGTTCCAGGGGTTTCCCTTTCCACTCATCGGAGATTCAGAGGGATGAGCTGGCACCAG
CTGGGACAGGGGTGTCCCGTGAGGCTGTGTCGGGTCTGC

Figure 6 (continued)

>BAIP1
CGGCCGCGGACGAAGATGGCGACCGCCATGTACTTGGAGCACTATCTGGACAGTATCGAGAACCTT
CCCTGCGAACTTCAGAGGAACTTCCAGCTGATGCGAGAGCTGGACCAGAGGACGGAAGATAAGAAA
GCAGAGATTGACATCCTGGCTGCAGAGTACATCTCCACGGTGAAGACGCTGTCTCCAGACCAGCGC
GTGGAGCGCCTGCAGAAGATCCAGAACGCCTACAGCAAGTGCAAGGAATACAGTGACGACAAAGTG
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GAAGACACACCAAAGAAAAAGAAGCACAAAGGAGGGTCTGAGTTCACTGACACCATCCTGTCCGTG
CACCCCTCTGATGTGCTGGACATGCCCGTGGACCCAAACGAACCCACGTACTGCCTGTGCCACCAG
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>BAIP2
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TGCACCGCCATCCTCAAGTTTGCAGACGCCTTCGACCATCACAAGCTTCGATCTCAGGCCCAGTCC
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GTCTTCAAGTGCGTGCGCTGGATGCACTTCACTGAAGAAGATCAGGACTACTTAGAAGGGCTGCTG
ACCAAGCCCATCGTGAAGAAGTACTGCCTGGACGTTATTGAAGGGGCCCTGCAGATGCGCTATGGT
GACCTGTTGTACAAGTCTCTGGTGCCAGTGCCAAACAGCAGCAGCAGCAGTAGCAGCAGCAACTCT
CTTGTATCTGCAGCAGAAAATCCACCCCAGAGACTGGGTATGTGTGCCAAGGAGATGGTGATCTTC
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CATGACATCTATCTAGCTGCTCAGCCCAGGAAAGACCTCTGGGTGTATAAACCAGCTCAGAATAGT
TGGCAGCAACTTGCAGATCGCTTGCTGTGTCGTGAGGGCATGGATGTGGCATATCTCAATGGCTAC
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GTGGTTCAGAACTATCTTTATGCTGTCAACAGTAAGCGCATGCTTTGCTATGATCCTAGCCACAAT
ATGTGGCTGAACTGTGCTTCTCTTAAACGTAGTGACTTTCAGGAAGCATGTGTCTTCAATGATGAA
ATCTATTGTATCTGTGACATCCCAGTCATGAAGGTCTACAACCCAGCTAGGGGAGAATGGAGGCGG
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GATGCACGGAGTGAGTCTAGTACTGAATGGGACTTAGATGGATTCAGTGAGCTGGACTCTGAGTCA
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GATCAGCAGGGTTCTTTG

Figure 6 (continued)

>BAIP3
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CAAGAACCAAAGATCCATAATGAGATGGCATCAACATCAGATAAAGGTGCCCAAGGAAGAAATGAC
AAGAAAGATTCTCAAGGAAGAAGTAATAAGGCATTACATCTGAAGAGTGATGCTGAATTTAAAAAG
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GAAGGTTTCGATTCCTTTAGCAGTTTGGTAAAGAGTGGTACTTACAAAGAGACAGAGTTTATGGTG
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AGTCACAGCAAAACCAGACAAGAAAAGAGAACTGAGATGGAATACTATACCCATGAGAAGCAAGAG
AAAGGCACTTTGAATTCAAATGCAGCTTATGAACAAAGTCATTTCTTCAATAAAAATTATACCGAA
GATATTTTCCCAGTGACACCACCGGAGTTAGAAGAAACCATTCGAGATGAAAAAATAAGAAGACTT
AAGCAGGTGCTGAGAGAGAAAGAAGCAGCTCTTGAAGAAATGCGTAAGAAGATGCACCAAAAA
>BARD1
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GATTTGAAAGAAGATAAACCTAGGAAAAGTTTGTTTAATGATGCAGGAAACAAGAAGAATTCAATT
AAAATGTGGTTTAGCCCTCGAAGTAAGAAAGTCAGATATGTTGTGAGTAAAGCTTCAGTGCAAACC
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CCTGCAGATGTTTCTGAGAGGGCTAAAAAGGCTTCTGCAAGATCTGGAAAAAAGCAAAAAAAGAAA
ACTTTAGCTGAAATCAACCAAAAATGGAATTTAGAGGCAGAAAAAGAAGATGGTGAATTTGACTCC
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GTGACTCCTGAGAAGGTCTGCAAAAATTATCTTACATCTAAGAAATCTTTGCCATTAGAAAATAAT
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>CA150
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CCTGCTCCTACAGCCACACCTGTGCAAACCGTTCCCCAGCCGCACCCTCAGACGTTACCTCCTGCT
GTTCCTCATTCAGTACCTCAGCCAACAACAGCAATACCTGCTTTTCCACCAGTAATGGTACCTCCG
TTTCGTGTTCCCCTTCCTGGCATGCCAATTCCACTTCCAGGTGTATTGCCAGGAATGGCCCCTCCT
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GCAGTTTCTGAATGGACTGAATATAAAACAGCAGATGGGAAGACATATTATTATAATAATAGAACA
TTAGAATCAACCTGGGAAAAACCCCAAGAACTAAAGGAAAAAGAAAAGTTAGAAGAGAAGATTAAA
GAGCCAATTAAAGAACCCTCTGAAGAGCCTCTGCCAATGGAGACGGAGGAGGAGGATCCTAAAGAA
GAGCCTATAAAGGAGATAAAGGAGGAGCCCAAAGAAGAGGAGATGACTGAAGAAGAAAAGGCTGCC
CAGAAGGCAAAGCCAGTTGCTACTGCTCCTATTCCTGGTACTCCATGGTGTGTCGTTTGGACTGGT
GATGAGCGGGTCTTCTTTTATAATCCCACCACTCGTCTTTCTATGTGGGACCGACCTGATGATCTG
ATTGGCAGGGCAGATGTTGACAAAATTATTCAGGAGCCCCCTCATAAAAAAGGAATGGAGGAATTG
AAGAAACTAAGGCACCCAACTCCGACAATGCTGTCGATCCAAAAGTGGCAATTCTCTATGAGTGCA
ATTAAAGAGGAACAAGAATTAATGGAAGAAATTAATGAAGATGAGCCTGTTAAAGCAAAAAAACGG
AAG

Figure 6 (continued)

>CGI-125
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CGGTTTCAGTTGGAGTTGGAATTTGTGCAATGTTTAGCCAACCCAAATTACCTTAATTTTCTTGCC
CAAAGAGGTTACTTCAAAGACAAAGCTTTTGTTAATTATCTTAAATACTTGCTTTACTGGAAAGAC
CCAGAATATGCCAAGTATCTAAAGTACCCTCAGTGTTTACACATGTTAGAGCTGCTCCAATATGAA
CACTTCCGAAAGGAGCTGGTGAATGCTCAGTGTGCGAAATTTATTGATGAACAGCAGATTCTACAT
TGGCAGCACTATTCCCGGAAGCGGATGCGCCTTCAGCAAGCCTTGGCAGAGCAGCAACAGCAAAAT
AACACATCGGGAAAA
>CGI-74
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AGTTTTCAGCAGCAGAAACTTCGAGTCTGTGAAGTCTGCTCTGCCTATTTAGGACTTCATGATAAT
GACAGACGACTGGCTGATCATTTTGGGGGTAAACTGCACCTGGGATTTATTGAAATAAGAGAGAAG
CTTGAAGAATTAAAGAGAGTCGTAGCTGAGAAGCAGGAGAAAAGAAACCAGGAACGGCTGAAACGA
AGAGAAGAGAGAGAGAGAAGAAAGGGAGAAGCTGAGGAGGTCCCGATCACACAGCAAGAATCCA
AAAAGG
>CLH-17
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GCAAACATTGGCTTCAGTACCCTGACTATGGAGTCTGACAAATTCATCTGCATTAGAGAAAAAGTA
GGAGAGCAGGCCCAGGTGGTAATCATTGATATGAATGACCCAAGTAATCCAATTCGAAGACCAATT
TCAGCAGACAGCGCCATCATGAATCCAGCTAGCAAAGTAATTGCACTGAAAGCTGGGAAAACTCTT
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CAGATCAGTGAAAAGCATGATGTGGTGTTCTTGATAACCAAGTATGGTTATATCCACCTCTATGAT
CTTGAGACT
>CLK1
GACGCGTGGGTTCTGGAACATCTGAATACAACAGACCCCAACAGTACTTTCCGCTGTGTCCAGATG
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GACTTCATTAAAGAAAATGGTTTTCTACCATTTCGACTGGATCATATCAGAAAGATGGCATATCAG
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CAGAAAACCAGGAAACGTAAATATTTTCACCACGATCGATTAGACTGGGATGAACACAGTTCTGCC
GGCAGATATGTTTCAAGACGCTGTAAACCTCTGAAGGAATTTATGCTTTCTCAAGATGTTGAACAT
GAGCGTCTCTTTGACCTCATTCAGAAAATGTTGGAGTATGATCCAGCCAAAAGAATTACTCTCAGA
GAAGCCTTAAAGCATCCTTTCTTTGACCTTCTGAAGAAAAGTATA

Figure 6 (continued)

>DRP-1
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ATCTTCGAGGGTATGGAGTGCCACGGCTCCCCACTAGTGGTCATCAGCCAGGGCAAGATCGTCTTT
GAAGACGGAAACATCAACGTCAACAAGGGCATGGGCCGCTTCATTCCGCGGAAGGCGTTCCCGGAG
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TATGACGGTCCTGTGTACGAGGTACCAGCTACACCCAAATATGCAACTCCCGCTCCTTCAGCCAAA
TCTTCGCCTTCTAAACACCAGCCCCCACCCATCAGAAACCTCCACCAGTCCAACTTCAGCTTATCA
GGTGCCCAGATAGATGACAACAATCCCAGGCGCACCGGCCACCGCATCGTGGCGCCCCCTGGTGGC
CGCTCCAACATCACCAGCCTCGGT
>EF1A
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GTTCGTGATATGAGACAGACAGTTGCGGTGGGTGTCATCAAAGCAGTGGACAAGAAGGCTGCTGGA
GCTGGCAAGGTCACCAAGTCTGCCCAGAAAGCTCAGAAGGCTAAA
>EF1G(bait)
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ACCTGGGTGTTCCCCACCTTGGGCATCATGCACCACAACAAACAGGCCACTGAGAATGCAAAGGAG
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CCCCAGGCTGAGCGGAAGGAGGAGAAAAAGGCGGCTGCCCCTGCTCCTGAGGAGGAGATGGATGAA
TGTGAGCAGGCGCTGGCTGCTGAGCCCAAGGCCAAGGACCCCTTCGCTCACCTGCCCAAGAGTACC
TTTGTGTTGGATGAATTTAAGCGCAAGTACTCCAATGAGGACACACTCTCTGTGGCACTGCCATAT
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CTCACTCAGACCTTCATGAGCTGCAATCTCATCACTGGAATGTTCCAGCGACTGGACAAGCTGAGG
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GTCTTCCGAGGCCAGGAGCTTGCCTTTCCGCTGAGTCCAGATTGGCAGGTGGACTACGAGTCATAC
ACATGGCGGAAACTGGATCCTGGCAGCGAGGAGACCCAGACGCTGGTTCGAGAGTACTTTTCCTGG
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Figure 6 (continued)

>EF1G(prey)
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ACCCCTGAATTTCTCCGCAAATTTCCTGCCGGCAAGGTCCCAGCATTTGAGGGTGATGATGGATTC
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ACCTGGGTGTTCCCCACCTTGGGCATCATGCACCACAACAAACAGGCCACTGAGAATGCAAAGGAG
GAAGTGAGGCGAATTCTGGGGCTGCTGGATGCTTACTTGAAGACGAGGACTTTTCTGGTGGGCGAA
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CCCCAGGCTGAGCGGAAGGAGGAGAAAAAGGCGGCTGCCCCTGCTCCTGAGGAGGAGATGGATGAA
TGTGAGCAGGCGCTGGCTGCTGAGCCCAAGGCCAAGGACCCCTTCGCTCACCTGCCCAAGAGTACC
TTTGTGTTGGATGAATTTAAGCGCAAGTACTCCAATGAGGACACACTCTCTGTGGCACTGCCATAT
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CTCACTCAGACCTTCATGAGCTGCAATCTCATCACTGGAATGTTCCAGCGACTGGACAAGCTGAGG
AAGAATGCCTTCGCCAGTGTCATCCTTTTTGGAACCAACAATAGCAGCTCCATTTCTGGAGTCTGG
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>FEZ1
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>G45IP1
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Figure 6 (continued)

>G45IP2
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CTTGGGGGCCCATCCTTCTTCAAAATCATCACGGCCAAGGCTGTCCTGAAGCTGCAGGCCGGAAAC
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CGGGAGAATGGCTTCCTGCTGCAGTACCTGGTGGCTATCCCCATGGAGAAAGGCCTTGACTCCCAA
GGCTGCTTCTGCGCAGGCTGCTCCCGGCAGATCGGCTTCTCCTTTGTACGACCCAAGCTCTGTGCC
TTCTCTGGCCTCTATTACTGTGACATCTGCCACCAAGACGATGCCTCAGTGATTCCGGCCAGGATC
ATCCACAACTGGGACCTCACCAAGCGCCCGATCTGCAGGCAGGCCCTGAAGTTTCTGACACAGATC
CGGGCCCAGCCCCTCATCAACCTGCAGATGGTGAACGCGTCTCTGTACGAGCATGTGGAGCGGATG
CACCTCATTGGGAGGAGACGGGAGCAGCTGAAGCTCCTGGGGGGATTACCTGGGCCTGTGCCGGAGT
GGCGCCCTGAAGGAGCTCAGCAAGAGGCTCAACCACAGGAATTATCTCTTGGAATCTCCGCATAGG
TTCAGTGTTGCTGACCTCCAACAGATCGCAGACGGGGTGTATGAAGGATTCCTCAAGGCCCTGATT
GAATTTGCCTCCCAGCATGTCTACCACTGCGACCTGTGCACCCAGCGCGGCTTCATCTGCCAGATC
TGCCAGCACCACGACATCATCTTCCCCTTTGAGTTTGACACCACAGTCAGGTGTGCCGAGTGCAAG
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AAGTACCAGGAACAGAACATTTTCGCC
>G45IP3
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AACACGAGAAAGCACATACCAAGGCCACCACACACCACCTGTCCAAAAAGGCCTTCGATACGGGAT
AATCCTATTTATTACCTCAGAAGTTTTTTTCTTCGCAGGATTTTTCTGAGCCTTTTACCACTCCAG
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GCTGCGGGTGAGGAGGACGAGGGCGACATCGCGCTGCAGATCCATTTTACGCTGATCCAGGCTTTC
TGCTGCGAGAACGACATCGACATAGTGCGCGTGGGCGATGTGCAGCGGCTGGCGGCTATCGTGGGC
GCCGGCGAGGAGGCGGGTGCGCCGGGCGACCTGCACTGCATCCTCATTTCGAACCCCAACGAGGAC
GCCTGGAAGGATCCCGCCTTGGAGAAGCTCAGCCTGTTTTGCGAGGAGAGCCGCAGCGTTAACGAC
TGGGTGCCCAGCATCACCCTCCCCGAG

Figure 6 (continued)

>GIT1
CCACAGATGGCTGACAGATCTCGGCAAAAGTGCATGTCTCAGAGCCTTGACTTATCCGAATTGGCC
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GTGACAGAGCGCAGTGCCGTGCCCTTCCTGCCTGTTAACCCGGAATACTCAGCCACGCGGAATCAG
GGGCGACAAAAGCTGGCCCGCTTTAATGCCCGAGAGTTTGCCACCTTGATCATCGACATTCTCAGT
GAGGCCAAGCGGAGACAGCAGGGCAAGAGCCTGAGCAGCCCCACAGACAACCTCGAGCTGTCTCTG
CGGAGCCAGAGTGACCTCGACGACCAACACGACTACGACAGCGTGGCCTCTGACGAGGACACAGAC
CAGGAGCCCCTGCGCAGCACCGGCGCCACTCGGAGCAACCGGGCCCGGAGCATGGACTCCTCGGAC
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GAGATCCACAAGCTGCAGGCGGAGAACCTGCAGCTCCGGCAGCCTCCAGGGCCGGTGCCCACACCT
CCACTCCCCAGTGAACGGGCGGAACACACACCCATGGCGCCAGGCGGGAGCACACACCGCAGGGAT
CGCCAGGCCTTTTCCATGTATGAACCTGGCTCTGCCCTGAAGCCCTTTGGGGGCCCCCCTGGGGAC
GAGCTCACTACGCGGCTGCAGCCTTTCCACAGCACTGAGCTAGAGGACGACGCCATCTATTCAGTG
CACGTCCCTGCTGGCCTTTACCGGATCCGGAAAGGGGTGTCTGCCTCAGCTGTGCCCTTCACTCCC
TCCTCCCCGCTGCTGTCCTGCTCCCAGGAGGGAAGCCGCCACACGAGCAAGCTTTCCCGCCACGGC
AGTGGAGCCGACAGTGACTATGAGAACACGCAAAGTGGGGACCCACTGCTGGGGCTGGAAGGGAAG
AGGTTTCTAGAGCTGGGCAAAGAGGAAGACTTCCACCCAGAGCTGGAAAGCCTGGATGGAGACCTA
GATCCTGGGCTTCCCAGCACAGAGGATGTCATCTTGAAGACAGAGCAGGTCACCAAGAACATTCAG
GAACTGTTGCGGGCAGCCCAGGAGTTCAAGCATGACAGCTTCGTGCCCTGCTCAGAGAAGATCCAT
TTGGCTGTGACCGAGATGGCCTCCCTCTTCCCAAAGAGGCCAGCCCTGGAGCCAGTGCGGAGCTCA
CTGCGGCTGCTCAACGCCAGCGCCTACCGGCTGCAGAGTGAGTGCCGGAAGACAGTGCCCCCAGAG
CCCGGCGCCCCAGTGGACTTCCAGCTGCTGACTCAGCAGGTGATCCAGTGCGCCTATGACATCGCC
AAGGCTGCCAAGCAGCTGGTCACCATCACCACCCGAGAGAAGAAGCAG
>hADA3
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GATTCTCCTATTCCTGACATGTCTGGGAAAGAATCAGGGGCTGACGGGGCAAGCACCTCCCCTCGC
AATCAGAACAAGCCCTTCAGTGTGCCGCATACTAAGTCCCTGGAGAGCCGCATCAAGGAGGAGCTA
ATTGCCCAGGGCCTTTTGGAGTCTGAGGACCGCCCCGCAGAGGACTCCGAGGATGAGGTCCTTGCT
GAGCTTCGCAAACGGCAGGCTGAGCTGAAGGCACTTAGTGCCCACAACCGCACCAAGAAGCACGAC
CTGCTGAGGCTGGCAAAGGAGGAGGTGAGCCGGCAGGAGCTGAGGCAGCGGGTGCGCATGGCTGAC
AACGAGGTCATGGACGCCTTTCGCAAGATCATGGCTGCCCGGCAGAAGAAGCGGACTCCCACCAAG
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Figure 6 (continued)

>HBO1
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CCGCCGGCAATGCCGCGAAGGAAGAGGAATGCAGGCAGTAGTTCAGATGGAACCGAAGATTCCGAT
TTTTCTACAGATCTCGAGCACACAGACAGTTCAGAAAGTGATGGCACATCCCGACGATCTGCTCGA
GTCACCCGCTCCTCAGCCAGGCTAAGCCAGAGTTCTCAAGATTCCAGTCCTGTTCGAAATCTGCAG
TCTTTTGGCACTGAGGAGCCTGCTTACTCTACCAGAAGAGTGACCCGTAGTCAGCAGCAGCCTACC
CCAGTGACACCGAAAAAATACCCTCTTCGGCAGACTCGTTCATCTGGTTCAGAAACTGAGCAAGTG
GTTGATTTTTCAGATAGAGAAACTAAAAATACAGCTGATCATGATGAGTCACCGCCTCGAACTCCA
ACTGGAAATGCGCCTTCTTCTGAGTCTGACATAGACATCTCCAGCCCCAATGTATCTCACGATGAG
AGCATTGCCAAGGACATGTCCCTGAAGGACTCAGGCAGTGATCTCTCTCATCGCCCCAAGCGCCGT
CGCTTCCATGAAAGCTACAACTTCAATATGAAGTGTCCTACACCAGGCTGTAACTCTCTAGGACAC
CTTACAGGAAAACATGAGAGACATTTCTCCATCTCAGGATGCCCACTGTATCATAACCTCTCAGCT
GACGAATGCAAGGTGAGAGCACAGAGCCGGGATAAGCAGATAGAAGAAAGGATGCTGTCTCACAGG
CAAGATGACAACAACAGGCATGCAACCAGGCACCAGGCACCAACGGAGAGGCAGCTTCGATATAAG
GAAAAAGTGGCTGAACTCAGGAAGAAAAGAAATTCTGGACTGAGCAAAGAACAGAAAGAGAAATAT
ATGGAACACAGACAGACCTATGGGAACACACGGGAACCTCTTTTAGAAAACCTGACAAGCGAGTAT
GACTTGGATCTTTTCCGAAGAGCACAAGCCCGGGCTTCAGAGGATTTGGAGAAGTTAAGGCTGCAA
GGCCAAATCACAGAGGGAAGCAACATGATTAAAACAATTGCTTTTGGCCGCTATGAGCTTGATACC
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GTGGAGCCCTTCCTGTTCTATGTTATGACAGAGGCGGACAACACTGGCTGTCACCTGATTGGATAT
TTTTCTAAGGAAAAGAATTCATTCCTCAACTACAACGTCTCCTGTATCCTTACTATGCCTCAGTAC
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GTACTTCTCCGCTACCTGCATAATTTTCAAGGCAAAGAGATTTCTATCAAAGAAATCAGTCAGGAG
ACGGCTGTGAATCCTGTGGACATTGTCAGCACTCTGCAAGCCCTTCAGATGCTCAAATACTGGAAG
GGAAAACACCTAGTTTTAAAGAGACAGGACCTGATTGATGAGTGGATAGCCAAAGAGGCCAAAAGG
TCCAACTCCAATAAAACCATGGATCCCAGCTGCTTAAAATGGACCCCTCCCAAGGGCACT

Figure 6 (continued)

>HD1.7
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CAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAACAGCCGCCACCGCCG
CCGCCGCCGCCGCCGCCTCCTCAGCTTCCTCAGCCGCCGCCGCAGGCACAGCCGCTGCTGCCTCAG
CCGCAGCCGCCCCCGCCGCCGCCCCCGCCGCCACCCGGCCCGGCTGTGGCTGAGGAGCCGCTGCAC
CGACCAAAGAAAGAACTTTCAGCTACCAAGAAAGACCGTGTGAATCATTGTCTGACAATATGTGAA
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CTTTTTCTGCTGTGCAGTGATGACGCAGAGTCAGATGTCAGGATGGTGGCTGACGAATGCCTCAAC
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>HDd1.0
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CCGCAGCCGCCCCCGCCGCCGCCCCCGCCGCCACCCGGCCCGGCTGTGGCTGAGGAGCCGCTGCAC
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AATTTTGCAAATGACAATGAAATTAAGGTTTTGTTAAAGGCCTTCATAGCGAACCTGAAGTCAAGC
TCCCCCACCATTCGGCGGACAGCGGCTGGATCAGCAGTGAGCATCTGCCAGCACTCAAGAAGGACA
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Figure 6 (continued)

>HDd1.3
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GCAGCTGTTCCCAAAATTATGGCTTCTTTTGGCAATTTTGCAAATGACAATGAAATTAAGGTTTTG
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GCAGTGAGCATCTGCCAGCACTCAAGAAGGACACAATATTTCTATAGTTGGCTACTAAATGTGCTC
TTAGGCTTACTCGTTCCTGTCGAGGATGAACACTCCACTCTGCTGATTCTTGGCGTGCTGCTCACC
CTGAGGTATTTGGTGCCCTTGCTGCAGCAGCAGGTCAAGGACACAAGCCTGAAAGGCAGCTTCGGA
GTGACAAGGAAAGAAATGGAAGTCTCTCCTTCTGCAGAGCAGCTTGTCCAGGTTTATGAACTGACG
TTACATCATACACAGCACCAAGACCACAATGTTGTGACCGGAGCCCTGGAGCTGTTGCAGCAGCTC
TTCAGAACGCCTCCACCCGAGCTTCTGCAAACCCTGACCGCAGTCGGGGGCATTGGGCAGCTCACC
GCTGCTAAGGAGGAGTCTGGTGGCCGAAGCCGTAGTGGGAGTATTGTGGAACTTATAGCTGGAGGG
GGTTCCTCATGCAGCCCTGTCCTTTCAAGAAAACAAAAAGGCAAAGTGCTCTTAGGAGAAGAAGAA
GCCTTGGAGGATGACTCTGAATCGAGATCGGATGTCAGCAGCTCTGCCTTAACAGCCTCAGTGAAG
GATGAGATCAGTGGAGAGCTGGCTGCTTCTTCAGGGGTTTCCACTCCAGGGTCAGCAGGTCATGAC
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>HDexQ20
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CCG
>HDexQ51
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>HIP1
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Figure 6 (continued)

>HIP11
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>HIP13
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CGCCGGGGAACCATTGGAGCTGGTCCCATCCCCATCAAGACACCCGTGATCCCTGTCAAGACCCCA
ACCGTCCCAGACCTCCCAGGGGTGTTGCCAGCCCCTCCAGATGGGCCAGAAGAGCGGGGGGAGCAC
AGCCCTGAGTCGCCATCTGTGGGTGAGGGCCCCCAAGGTGTCACCAGCATGCCCTCCTCAATGTGG
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CAGGCAATTCCAGAAAGTGAAGCTGAAGACCAGGAACGGGAACCCCCAAGTGCCACTGTCTCCCCA
GGCCAGATTCCAGAGAGTGACCCTGCAGACCTGAGCCCAAGGGATACTCCACAAGGAGAAGACATG
CTGAACGCCATCCGAAGGGGCGTGAAACTGAAGAAGACCACGACAAACGATCGCTCAGCCCCTCGC
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>HIP15
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ACAGCCAATGCGAGAACGAAGTTTCACGTTCTGAAAATGCTATTGAATTTGTCTGAAAATCCTGCT
GTGGCAAAAAAACTATTCAGTGCCAAAGCTCTTTCAATATTTGTGGGTCTCTTTAACATAGAAGAG
ACAAATGATAATATTCAAATTGTTATTAAAATGTTTCAGAATATCAGTAACATTATAAAAAGTGGA
AAGATGTCCTTAATTGATGATGATTTCAGTCTTGAGCCGCTTATTTCTGCATTTCGTGAATTTGAG
GAGTTAGCTAAGCAACTACAAGCCCAAATAGACAACCAAAATGATCCTGAGGTGGGACAACAAAGT
>HIP16
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GATTTCTTGAAAGAGAAGAGGGAAGCTGTGGAGGCGAGTAAGCCAAAGGACGTGGACCTGACACTA
CCTGGCTGGGGCGAGTGGGGTGGTGTGGGCCTAAAGCCCAGTGCCAAGAAAAGACGCCGGTTTCTC
ATTAAAGCCCCTGAGGGTCCTCCAAGAAAAGATAAGAATTTGCCAAATGTGATTATCAATGAGAAG
CGCAACATCCACGCAGCTGCTCATCAGGTACGAGTGCTTCCATATCCATTTACCCACCATTGGCAA
TTTGAAAGGACCATCCAGACCCCCATAGGATCCACATGGAACACCCAGAGGGCTTTCCAAAAGCTG
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CACAAAAAACAGCTGAAGAAATGCTCTGTAGAT
>HIP2
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GCACCAGTTTCTAGTCCAGAATACACCAAAAAAATAGAAAACCTATGTGCTATGGGCTTTGATAGG
AATGCAGTAATAGTGGCCTTGTCTTCAAAATCATGGGATGTAGAGACTGCAACAGAATTGCTTCTG
AGTAACTG

Figure 6 (continued)

>HIP5(bait)
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GAAAAAGGTGCAGAAATTCCAAAGACTATTAAAAAACTGAGGTGGTTTGATGAAACTAGCAATATA
GAAAACAATGCTGAAAACAGTCATTCACTGAAGAATAAAACAGGAACAACTCAACAGCATTCTCAA
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GCTGATACAAAGAAGTCCAGGGAGGATTCTATCTCTGAAAATGTTACGACTTTAGGAGGATCTGGA
GCAGACCATATGCCTTTGAACTGTTTTATACCTTCAGGTTATAACTTTGCTAAACATGCCTGGCCA
GCCTCAAAAAAAGAAGAAAGTAAAATCCCTGTACATGATGATTCTAAAACTAAGCAAGGTAAGCCA
CAAAGAGGTAGAGCAAAAATAATTAGAAAACCAGGATCTGCAAAAGTCCAATCAGGCTTTATATGT
ACAAACAGAAAAGGCGCTGTCATTCAACCACAGTCTGCAAGCAAAGTCAACATATTTACACAAGCT
CAGGGAAAATTAATTATACCTTGTCCTCCTCCTCAATCTACATCAAATATTAGAAGTGGTAAAAAT
ATACAAGTGTCTCAGTGTCAACCAGTAACTCCTGAAAATCCTCAAAACATTATTACACATAACTCT
TTTAATTCAAAACATGTGCTTCCAACAGAACACAGTTTGAATCAGTGGAATCAGGAAAGTAGTTCT
CCACTCTCAAATGCTTGTTCTGACCTAGTCACTGTGATACCATCACTGCCATCATATTGTTCTTCA
GAGTGCCAAACTTTCGCAAAAATAAATCATTCAAATGGCACTCAAGCAGTTGCCCGGCAAGATGCG
ACATTATATTGCACCCAAAGAAGTCCTGTTTGTGAAGAAAGTTATCCGTCTGTGACTCTAAGAACT
GCTGAAGAAGAATCAGTTCCCTTATGGAAAAGAGGTCCTAATGTCCTGCATCAAAATAAGAGGGCT
ACAGGGTCTACTGTTATGAGAAGAAAACGAATTGCTGAAACTAAGCGGAGAAATATTTTAGAGCAG
AAAAGACAAAACCCTGGATCTGTAGGACAGAAGTACAGTGAGCAAATTAATAATTTTGGACAAAGT
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CTTCTGGAGAAGAGAGAAGATAGAACCAGCAGCTGCAGAGACAAGAGA
>HIP5(prey)
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CTTCTGGAGAAGAGAGAAGATAGAACCAGCAGCTGCAGAGACAAGAGA

Figure 6 (continued)

>HMP
CAAGAACAAGTTAAAATTGAGTCTCTAGCCAAGAGCTTAGAAGATGCTCTGAGGCAAACTGCAAGT
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TTGAAAGCCGCCATGGACAATTCTGAGATTGCAGGCGAGAAGAAATCTGCTCAGTGGCGCACAGTG
GAGGGTGCATTGAAGGAACGCAGAAAGGCAGTAGATGAAGCTGCCGATGCCCTTCTCAAAGCCAAA
GAAGAGTTAGAGAAGATGAAAAGTGTGATTGAAAATGCAAAGAAAAAAGAGGTTGCTGGGGCCAAG
CCTCATATAACTGCTGCAGAGGGTAAACTTCACAACATGATAGTTGATCTGGATAATGTGGTCAAA
AAGGTCCAAGCAGCTCAGTCTGAGGCTAAGGTTGTATCTCAGTATCATGAGCTGGTGGTCCAAGCT
CGGGATGACTTTAAACGAGAGCTGGACAGTATTACTCCAGAAGTCCTTCCTGGGTGGAAAGGAATG
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>HP28
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AGCACCCCTAGCACCAGGATGGACGTGGTGCACCTCCAGGAGCAGTTAGACTTAAAGCTGCAGCAG
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GCACTGCAGGCTGAGCAGGGGAAGTCAGACATGGAGAGGAAAATCGCAGAATTGGAGACGGAAAAG
AGAGACCTGGAGAGGCAAGTGAACGAGCAGAAGGCAAAATGTGAAGCCACTGAGAAGCGGGAGAGC
GAGAGGCGGCAGGTGGAGGAGAAGAAGCACAATGAGGAGATTCAGTTCCTGAAGCGAACAAATCAG
CAGCTGAAGGCCCAACTGGAAGGCATTATTGCACCAAAGAAG

Figure 6 (continued)

>HSPC232
CGGCGGCGAGCGGACGGCTGCATTTACGGGGTCTCCCGGAGGGCCAGAGTCGTGGCTTACAGAAGA
GACGAAATGTGGTCTGAGGGACGATATGAATATGAAAGAATTCCGAGAGAACGAGCACCTCCTCGA
AGTCATCCCAGTGATGAATCTGGTTATAGATGGACAAGAGACGATCATTCTGCAAGCAGGCAACCT
GAATACAGGGACATGAGAGATGGCTTTAGAAGAAAAAGTTTCTACTCTTCCCATTATGCGAGAGAG
CGGTCTCCTTATAAAAGGGACAATACTTTTTTCAGAGAATCACCTGTTGGCCGAAAGGATTCTCCA
CACAGCAGATCTGGTTCCAGTGTCAGTAGCAGAAGCTACTCTCCAGAAAGGAGCAAATCATACTCT
TTCCATCAGTCTCAACATAGAAATAAAGAGAGGCCTGTCCAGTCTTTGAAAACATCAAGAGATACT
TCACCCTCAAGTGGTTCAGCAGTTTCTTCATCAAAGGTGTTAGACAAACCCAGTAGGCTAACTGAA
AAGGAACTTGCTGAGGCTGCAAGCAAGTGGGCTGCTGAAAAGCTAGAGAAATCAGATGAAAGTAAC
TTGCCTGAAATTTCTGAGTATGAGGCGGGATCCACAGCACCATTGTTTACTGACCAGCCAGAGGAA
CCTGAGTCAAACACAACACATGGGATAGAATTATTTGAAGATAGTCAGCTAACCACTCGCTCTAAA
GCAATAGCATCAAAAACCAAAGAGATTGAACAGGTTTACCGACAAGACTGTGAAACTTTCGGGATG
GTGGTGAAAATGCTGATTGAAAAAGATCCTTCATTAGAAAAGTCTATACAGTTTGCATTGAGGCAG
AATTTACATGAAATAGGTGAGCGGTGTGTTGAAGAACTCAAGCATTTCATTGCAGAGTATGATACT
TCCACTCAAGATTTTGGAGAGCCTTTT
>HYPA
GGCCGCCGGCGGAGCAGTCTGAGCCCGACGATGAGGCCGGGGACGGGAGCTGAGCGTGGAGGCCTC
ATGATGGGGCACCCTGGCATGCATTATGCCCCAATGGGAATGCACCCTATGGGTCAGAGAGCGAAT
ATGCCTCCTGTACCTCATGGAATGATGCCGCAGATGATGCCCCCTATGGGAGGGCCACCAATGGGA
CAAATGCCTGGAATGATGTCGTCAGTAATGCCTGGAATGATGATGTCTCATATGTCTCAGGCTTCC
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AAATCAATGTGGACTGAACATAAATCACCTGATGGAAGGACTTACTACTACAACACTGAAACCAAA
CAGTCTACCTGGGAGAAACCAGATGATCTTAAAACACCTGCTGAGCAACTCTTATCTAAATGCCCC
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TGGGCCAAACCTAAAGAACTTGAGGATCTTGAAGGATACCAGAATACCATTGTTGCTGGAAGTCTT
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ACAACATCAACAGCTCCAGTCCCTACAACAGAAATTCCGACCACAATGAGCACCATGGCTGCTGCC
GAAGCAGCAGCTGCTGTTGTTGCAGCAGCAGCAGCGGCAGCAGCAGCAGCAGCTGCAGCCAATGCT
AATGCTTCCACTTCTGCTTCTAATACTGTCAGTGGAACTGTTCCAGTTGTTCCTGAGCCTGAAGTT
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CAACTTACTAGTACCCCTGCTATTCAGGATCAAAGTGTGGAAGTATCCAGTAATACTGGAGAAGAA
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AAAGAAAAAAAAAAAAAAAAAAAAAAA
>HZFH
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CTTCAGATGTCCGAGCGCAGCATCCTCAGCCGGCTGGCCAGCAAGGGCACGGAGCCTCACCCCACA
CCGGCCTACCCGCCGGGTCCCTACGCTACACCTCCGGGGTACGGGGCGGCCTTCAGCGCCGCACCC
GTAGGGGCCCTGGCCGCCGCAGGCGCCAATTACAGCCAGATGCCTGCAGGGTCCTTCATCACAGCC
GCCACCAACGGCCCTCCAGTGCTTGTGAAGAAGGAGAAGGAAATGGTGGGGGCATTGGTGTCAGAC
GGGCTGGATCGGAAGGAGCCCCGAGCCGGGGAGGTGATCTGTATAGACGAC

Figure 6 (continued)

>IKAP
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ACTGAAAACCTGAAAGATGAAGTATACCATATTTTAAAGGTACTCTTTCTCTTTGAGTTTGATGAA
CAAGGAAGGGAATTACAGAAGGCCTTTGAAGATACGCTGCAGTTGATGGAAAGGTCACTTCCAGAA
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AGTATCATGGCATCTTATCAGCAACAGAAGACTTCGGTTCCTGTTCTTGATGCTGAGCTTTTTATA
CCACCAAAGATCAACAGAAGAACCCAGTGGAAGCTGAGCCTGCTAGAC
>IMPD2
GACTTTCTCATTCTCCCTGGGTACATCGACTTCACTGCAGACCAGGTGGACCTGACTTCTGCTCTG
ACCAAGAAAATCACTCTTAAGACCCCACTGGTTTCCTCTCCCATGGACACAGTCACAGAGGCTGGG
ATGGCCATAGCAATGGCGCTTACAGGCGGTATTGGCTTCATCCACCACAACTGTACACCTGAATTC
CAGGCCAATGAAGTTCGGAAAGTGAAGAAATATGAACAGGGATTCATCACAGACCCTGTGGTCCTC
AGCCCCAAGGATCGCGTGCGGGATGTTTTTGAGGCCAAGGCCCGGCATGGTTTCTGCGGTATCCCA
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CTCAAAGAGGAGGAACATGACTGTTTCTTGGAAGAGATAATGACAAAGAGGGAAGACTTGGTGGTA
GCCCCTGCAGGCATCACACTGAAGGAGGCAAATGAAATTCTGCAGCGCAGCAAGAAGGGAAAGTTG
CCCATTGTAAATGAAGATGATGAGCTTGTGGCCATCATTGCCCGGACAGACCTGAAGAAGAATCGG
GACTACCCACTAGCCTCCAAAGATGCCAAGAAACAGCTGCTGTGTGGGGCAGCCATTGGCACTCAT
GAGGATGACAAGTATAGGCTGGACTTGCTCGCCCAGGCTGGTGTGGATGTAGTGGTTTTGGACTCT
TCCCAGGGAAATTCCATCTTCCAGATCAATATGATCAAGTACATCAAAGACAAATACCCTAATCTC
CAAGTCATTGGAGGCAATGTGGTCACTGCTGCCCAGGCCAAGAACCTCATTGATGCAGGTGTGGAT
GCCCTGCGGGTGGGCATGGGAAGTGGCTCCATCTGCATTACGCAGGAAGTGCTGGCCTGTGGGCGG
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>KPNA2
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GCTCTAGGAAACATTGCAGGTGATGGCTCAGTGTTCCGAGACTTGGTTATTAAGTACGGTGCAGTT
GACCCACTGTTGGCTCTCCTTGCAGTTCCTGATATGTCATCTTTAGCATGTGGCTACTTACGTAAT
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CAGATTCTTCCTACCTTAGTTCGGCTCCTGCATCATGATGATCCAGAAGTGTTAGCAGATACCTGC
TGGGCTATTTCCTACCTTACTGATGGTCCAAATGAACGAATTGGCATGGTGGTGAAAACAGGAGTT
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GACAAAATTGAAGCTCTACAAAACCATGAAAATGAGTCTGTGTATAAGGCTTCGTTAAGCTTAATT
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Figure 6 (continued)

>KPNB1
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GTTGTATTGAATACTCTTCAGCAGGCCTCCCAAGCCCAGGTGGACAAGTCAGACTATGACATGGTG
GATTATCTGAATGAGCTAAGGGAAAGCTGCTTGGAAGCCTATACTGGAATCGTCCAGGGATTAAAG
GGGGATCAGGAGAACGTACACCCGGATGTGATGCTGGTACAACCCAGAGTAGAATTTATTCTGTCT
TTCATTGACCACATTGCTGGAGATGAGGATCACACAGATGGAGTAGTAGCTTGTGCTGCTGGACTA
ATAGGGGACTTATGTACAGCATTTGGGAAGGATGTACTGAAATTAGTAGAAGCTAGGCCAATGATC
CATGAATTGTTAACTGAAGGGCGGAGATCGAAGACTAACAAAGCAAAAACCCTTGCTACATGGGCA
ACAAAAGAACTGAGGAAACTGAAGAACCAAGCT
>Ku70
AAGACCCGGACCTTTAATACAAGTACAGGCGGTTTGCTTCTGCCTAGCGATACCAAGAGGTCTCAG
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CCAGGTTTGATGCTCATGGGTTTCAAGCCGTTGGTACTGCTGAAGAAACACCATTACCTGAGGCCC
TCCCTGTTCGTGTACCCAGAGGAGTCGCTGGTGATTGGGAGCTCAACCCTGTTCAGTGCTCTGCTC
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TATTTTGTGGCTTTGGTGCCACAGGAAGAAGAGTTGGATGACCAGAAAATTCAGGTGACTCCTCCA
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ATGGCAACTCCAGAGCAGGTGGGCAAGATGAAGGCTATCGTTGAGAAGCTTCGCTTCACATACAGA
AGTGACAGCTTTGAGAACCCCGTGCTGCAGCAGCACTTCAGGAACCTGGAGGCCTTGGCCTTGGAT
TTGATGGAGCCGGAACAAGCAGTGGACCTGACATTGCCCAAGGTTGAAGCAATGAATAAAAGACTG
GGCTCCTTGGTGGATGAGTTTAAGGAGCTTGTTTACCCACCAGATTACAATCCTGAAGGGAAAGTT
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GAGCTGAAGACCCACATCAGCAAGGGTACGCTGGGCAAGTTCACTGTGCCCATGCTGAAAGAGGCC
TGCCGGGCTTACGGGCTGAAGAGTGGGCTGAAGAAGCAGGAGCTGCTGGAAGCCCTCACCAAGCAC
TTCCAGGAC
>LUC7B1
GTCGACGCGGTCGCGGTCGACGCGGCCGCGGTTTCTGCAAAGGCAGAAAAAGTACATGAGTTAAAT
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CAGAAGATTCTTATGGAAGTGGAAAAAGTTCGTGCGAAGAAAAAAGAAGCTGAGGAAGAATACAGA
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CTTGGTCTCCATGACAATGACCGTCGCCTGGCAGACCACTTCGGTGGCAAGTTACACTTGGGGTTC
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CAGGATCGCTTGAGGAGGAGAGAGGAGAGGGAACGGGAGGAGCGTCTGAGCAGGAGGTCGGGATCA
AGAACCAGAGATCGCAGGAGGTCACGCTCCCGGGATCGGCGTCGGAGGCGGTCAAGATCTACCTCC
CGAGAGCGACGGAAATTGTCCCGGTCCCGGTCCCGAGATAGACATCGGCGCCACCGCAGCCGTTCC
CGGAGCCACAGCCGGGGACATCGTCGGGCTTCCCGGGACCGAAGTGCGAAATACAAGTTCTCCAGA
GAGCGGGCATCCAGAGAGGAGTCCTGGGAGAGCGGGCGGAGCGAGCGAGGGCCCCCGGACTGGAGG
CTTGAGAGCTCCAACGGGAAGATGGCTTCACGGAGGTCAGAAGAGAAGGAGGCCGGCGAGATC
>MAGEH1
GCATCCTTCCCTAGGACTGCTGTAAGCTTTGAGCCTCTAGCAGGAGACATGCCTCGGGGACGAAAG
AGTCGGCGCCGCCGTAATGCGAGAGCCGCAGAAGAGAACCGCAACAATCGCAAAATCCAGGCCTCA
GAGGCCTCCGAGACCCCTATGGCCGCCTCTGTGGTAGCGAGCACCCCCGAAGACGACCTGAGCGGC
CCCGAGGAAGACCCGAGCACTCCAGAGGAGGCCTCTACCACCCCTGAAGAAGCCTCGAGCACTGCC
CAAGCACAAAAGCCTTCAGTGCCCCGGAGCAATTTTCAGGGCACCAAGAAAAGTCTCCTGATGTCT
ATATTAGCGCTCATCTTCATCATGGGCAACAGCGCCAAGGAAGCTCTGGTCTGGAAAGTGCTGGGG
AAGTTAGGAATGCAGCCTGGACGTCAGCACAGCATCTTTGGAGATCCGAAGAAGATCGTCACAGAA
GAGTTTGTGCGCAGAGGGTACCTGATTTATAAACCGGTGCCCCGTAGCAGTCCGGTGGAGTATGAG
TTCTTCTGGGGGCCCCGAGCACACGTGGAATCGAGCAAACTGAAAGTCATGCATTTTGTGGCAAGG
GTTCGTAACCGATGCTCTAAAGACTGGCCTTGTAATTATGACTGGGATTCGGACGATGATGCAGAG
GTTGAGGCTATCCTCAATTCAGGTGCTAGGGGTTATTCCGCCCCT

Figure 6 (continued)

>MAP1lc3
CAGCGGCGGAGCTTCGCCGACCGCTGTAAGGAGGTACAGCAGATCCGCGACCAGCACCCCAGCAAA
ATCCCGGTGATCATCGAGCGCTACAAGGGTGAGAAGCAGCTGCCCGTCCTGGACAAGACCAAGTTT
TTGGTCCCGGACCATGTCAACATGAGCGAGTTGGTCAAGATCATCCGGCGCCGCCTGCAGCTGAAC
CCCACGCAGGCCTTCTTCCTGCTGGTGAACCAGCACAGCATGGTGAGTGTGTCCACGCCCATCGCG
GACATCTACGAGCAGGAGAAAGACGAGGACGGCTTCCTCTATATGGTCTACGCCTCCCAGGAAACC
TTCGGCTTC
>mHAP1
CCGAAAGAGCAGGTGCAGAGCGGTGCGGGAGACGGGACAGGGTCGGGGGACCCAGCAGCAGGCACC
CCCACGACCCAGCCTGCAGTTGGTCCCGCTCCGGAGCCCTCGGCGGAGCCCAAACCTGCTCCAGCG
CAGGGAACCGGGTCCGGACAAAAATCAGGATCCCGAACCAAGACAGGAAGCTTTTGTCGGTCCATG
ATCATTGGTGATTCGGACGCACCATGGACCCGCTACGTATTCCAGGGGCCTTACGGTCCCCGGGCC
ACTGGCCTGGGCACTGGAAAGGCCGAGGGAATCTGGAAGACACCAGCCGCGTACATCGGCCGGAGG
CCCGGCGTGTCCGGCCCTGAGCGTGCGGCGTTTATTCGAGAGCTGCAGGAAGCGTTGTGTCCTAAT
CCACCACCCACGAAGAAGATCACCGAAGATGATGTCAAAGTGATGTTGTATTTGCTGGAAGAGAAA
GAACGGGACCTGAACACAGCCGCCCGGATCGGCCAGTCCCTGGTGAAACAGAACAGTGTCTTGATG
GAGGAGAATAATAAGCTGGAAACCATGCTGGGCTCAGCCAGGGAGGAGATTTTACATCTCCGGAAG
CAGGTGAACCTGCGAGATGACCTTCTTCAGCTCTACTCAGACTCTGATGACGATGATGATGAGGAA
GACGAGGAAGACGAGGAAGAGGGCGAAGAGGAGGAACGAGAAGGACAGAGGGATCAAGACCAGCAG
CACGACCACCCCTATGGTGCCCCCAAGCCACACCCTAAGGCTGAGACAGCGCACCGCTGCCCACAG
CTGGAAACCCTGCAGCAGAAGCTCAGGCTTCTGGAGGAAGAGAACGACCACCTGCGAGAGGAGGCC
TCCCACCTTGACAACCTGGAGGACGAAGAGCAGATGCTCATTCTGGAATGTGTGGAGCAGTTCTCT
GAAGCCAGCCAGCAGATGGCAGAGCTATCGGAAGTGCTGGTGTTGAGGCTGGAAGGCTATGAGAGG
CAGCAGAAAGAGATCACTCAGCTGCAGGCCGAGATCACCAAGCTACAACAGCGTTGTCAGTCTTAT
GGGGCCCAGACGGAGAAACTGCAGCAGATGCTGGCCTCAGAGAAGGGGATCCACTCGGAGAGCCTG
CGAGCTGGCTCCTACATGCAGGATTATGGGAGCAGGCCTCGTGACCGCCAGGAGGATGGGAAGAGT
CATCGCCAGCGCTCCTCCATGCCCGCAGGCTCTGTCACCCACTATGGATACAGTGTGCCTCTGGAT
GCACTTCCAAGTTTCCCAGAGACACTGGCTGAGGAGCTCCGAACATCTCTGAGGAAGTTCATCACT
GACCCTGCGTATTTCATGGAGAGACGTGACACTCACTGCAGGGAGGGGCGTAAGAAGGAGCAGAGG
GCGATGCCACCCCCACCGGCTCA
>mp53
GTCACCGAGACCCCTGGGCCAGTGGCCCCTGCCCCAGCCACTCCATGGCCCCTGTCATCTTTTGTC
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CCTGTGCAGTTGTGGGTCAGCGCCACACCTCCAGCTGGGAGCCGTGTCCGCGCCATGGCCATCTAC
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GATGGCCTGGCTCCTCCCCAGCATCTTATCCGGGTGGAAGGAAATTTGTATCCCGAGTATCTGGAA
GACAGGCAGACTTTTCGCCACAGCGTGGTGGTACCTTATGAGCCACCCGAGGCCGGCTCTGAGTAT
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TGCCCTGAACTGCCCCCAGGGAGCGCAAAGAGAGCGCTGCCCACCTGCACAAGCGCCTCTCCCCCG
CAAAAGAAAAAACCACTTGATGGAGAGTATTTCACCCTCAAGATCCGCGGGCGTAAACGCTTCGAG
ATGTTCCGGGAGCTGAATGAGGCCTTAGAGTTAAAGGATGCCCATGCTACAGAGGAGTCTGGAGAC
AGCAGGGCTCACTCCAGCTACCTGAAGACCAAGAAGGGCCAGTCTACTTCCCGCCATAAAAAAACA
ATGGTCAAGAAAGTGGGGCCTGACTCAGAC

Figure 6 (continued)

>NAG4
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GTGACTGATTTTATTGCTCCTGGCTACTCCATGATCATTAAACACCCAATGGATTTTAGTACCATG
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TGTACTAATGCCATGATTTACAATAAACCAGAGACCATTTATTATAAAGCTGCAAAGAAGCTGTTG
CACTCAGGAATGAAAATTCTTAGCCAGGAAAGAATTCAGAGCCTGAAGCAGAGCATAGACTTCATG
GCTGACTTGCAGAAAACTCGAAAGCAGAAAGATGGAACAGACACCTCACAGAGTGGGGAGGACGGA
GGCTGCTGGCAGAGAGAGAGAGAGGACTCTGGAGATGCCGAAGCACACGCCTTCAAGAGTCCCAGC
AAAGAAAATAAAAAGAAAGACAAAGATATGCTTGAAGATAAGTTTAAAAGCAATAATTTAGAGAGA
GAGCAGGAGCAGCTTGACCGCATCGTGAAGGAATCTGGAGGAAAGCTGACCAGGCGGCTTGTGAAC
AGTCAGTGCGAATTTGAAAGAAGAAAACCAGATGGAACAACGACGTTGGGACTTCTCCATCCTGTG
GATCCCATTGTAGGAGAGCCAGGCTACTGCCCTGTGAGACTGGGAATGACAACTGGAAGACTTCAG
TCTGGAGTGAATACTTTGCAGGGGTTCAAAGAGGATAAAAGGAACAAAGTCACTCCAGTGTTATAT
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GATGATTCTGATTTAATCTATTCAACCTATGGGGAAGACTCTGATCTTCCAAGTGATTTCAGCATC
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TCCAGTACTCAAGACAGGCTCATAGCGCTGAAAGCAGTAACAAATTTTGGCGTTCCAGTTGAAGTT
TTTGACTCTGAAGAAGCTGAAATATTCCAGAAGAAACTTGATGAGACCACCAGATTGCTCAGGGAA
CTCCAGGAAGCCCAGAATGAACGTTTGAGCACCAGACCCCCTCCGAACATGATCTGTCTCTTGGGT
CCCTCATACAGAGAAATGCATCTTGCTGAACAAGTGACCAATAATCTTAAAGAACTTGCACAGCAA
GTAACTCCAGGTGATATCGTAAGCACGTATGGAGTTCGAAAAGCAATGGGGATTTCCATTCCTTCC
CCCGTCATGGAAAACAACTTTGTGGATTTGACAGAAGACACTGAAGAACCTAAAAAGACGGATGTT
GCTGAGTGTGGACCTGGTGGAAGT

Figure 6 (continued)

>NEFL
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ACCATGAGTTCCTTCAGCTACGAGCCGTACTACTCGACCTCCTACAAGCGGCGCTACGTGGAGACG
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TCGGCGCCGGTGTCTTCCTCGCTGTCCGTGCGCCGCAGCTACTCCTCCAGCTCTGGATCGTTGATG
CCCAGTCTGGAGAACCTCGACCTGAGCCAGGTAGCCGCCATCAGCAACGACCTCAAGTCCATCCGC
ACGCAGGAGAAGGCGCAGCTCCAGGACCTCAATGACCGCTTCGCCAGCTTCATCGAGCGCGTGCAC
GAGCTGGAGCAGCAGAACAAGGTCCTGGAAGCCGAGCTGCTGGTGCTGCGCCAGAAGCACTCCGAG
CCATCCCGCTTCCGGGCGCTGTACGAGCAGGAGATCCGCGACCTGCGCCTGGCGGCGGAAGATGCC
ACCAACGAGAAGCAGGCGCTCCAGGGCGAGCGCGAAGGGCTGGAGGAGACCCTGCGCAACCTGCAG
GCGCGCTATGAAGAGGAGGTGCTGAGCCGCGAGGACGCCGAGGGCCGGCTGATGGAAGCGCGCAAA
GGCGCCGACGAGGCGGCGCTCGCTCGCGCCGAGCTCGAGAAGCGCATCGACAGCTTGATGGACGAA
ATCTCTTTTCTGAAGAAAGTGCACGAAGAGGAGATCGCCGAACTGCAGGCGCAGATCCAGTACGCG
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CAGTACGAGAAGCTGGCCGCCAAGAACATGCAGAACGCTGAGGAATGGTTCAAGAGCCGCTTCACC
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AGCCGTCGTCTGCTCAAGGCCAAGACCCTGGAAATCGAAGCATGCCGGGGCATGAATGAAGCGCTG
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GACCTCCTCAACGTGAAGATGGCTTTGGATATTGAGATTGCAGCTTACAGGAAACTCTTGGAAGGC
GAGGAGACCCGACTCAGTTTCACCAGCGTGGGAAGCATAACCAGTGGCTACTCCCAGAGCTCCCAG
GTCTTTGGCCGATCTGCCTACGGCGGTTTACAGACCAGCTCCTATCTGATGTCCACCCGCTCCTTC
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AAGAAAGTTGAAGGTGCTGGGGAGGAACAAGCAGCTAAGAAGAAAGAT
>p53
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CTGTCCCCGGACGATATTGAACAATGGTTCACTGAAGACCCAGGTCCAGATGAAGCTCCCAGAATG
CCAGAGGCTGCTCCCCCCGTGGCCCCTGCACCAGCAGCTCCTACACCGGCGGCCCCTGCACCAGCC
CCCTCCTGGCCCCTGTCATCTTCTGTCCCTTCCCAGAAAACCTACCAGGGCAGCTACGGTTTCCGT
CTGGGCTTCTTGCATTCTGGGACAGCCAAGTCTGTGACTTGCACGTACTCCCCTGCCCTCAACAAG
ATGTTTTGCCAACTGGCCAAGACCTGCCCTGTGCAGCTGTGGGTTGATTCCACACCCCCGCCCGGC
ACCCGCGTCCGCGCCATGGCCATCTACAAGCAGTCACAGCACATGACGGAGGTTGTGAGGCGCTGC
CCCCACCATGAGCGCTGCTCAGATAGCGATGGTCTGGCCCCTCCTCAGCATCTTATCCGAGTGGAA
GGAAATTTGCGTGTGGAGTATTTGGATGACAGAAACACTTTTCGACATAGTGTGGTGGTGCCCTAT
GAGCCGCCTGAGGTTGGCTCTGACTGTACCACCATCCACTACAACTACATGTGTAACAGTTCCTGC
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GAGAATCTCCGCAAGAAAGGGGAGCCTCACCACGAGCTGCCCCCAGGGAGCACTAAGCGAGCACTG
CCCAACAACACCAGCTCCTCTCCCCAGCCAAAGAAGAAACCACTGGATGGAGAATATTTCACCCTT
CAGATCCGTGGGCGTGAGCGCTTCGAGATGTTCCGAGAGCTGAATGAGGCCTTGGAACTCAAGGAT
GCCCAGGCTGGGAAGGAGCCAGGGGGGAGCAGGGCTCACTCCAGCCACCTGAAGTCCAAAAAGGGT
CAGTCTACCTCCCGCCATAAAAAACTCATGTTCAAGACAGAAGGGCCTGACTCAGAC

Figure 6 (continued)

>PFN2
GCTCCTCGCCGTCCGCGCTGCAGTGCGAAGGGCTCGAAGATGGCCGGTTGGCAGAGCTACGTGGAT
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TGGGCAGCCACGGCCGGGGGCGTCTTTCAGAGCATTACGCCAATAGAAATAGATATGATTGTAGGA
AAAGACCGGGAAGGTTTCTTTACCAACGGTTTGACTCTTGGCGCGAAGAAATGCTCAGTGATCAGA
GATAGTCTATACGTCGATGGTGACTGCACAATGGACATCCGGACAAAGAGTCAAGGTGGGCGAGCCA
ACATACAATGTGGCTGTCGGCAGAGCTGGTAGAGTCTTGGTCTTTGTAATGGGAAAAGAAGGGGTC
CATGGAGGCGGATTGAATAAGAAGGCATACTCAATGGCAAAATACTTGAGAGACTCTGGGTTC
>PIASy(bait)
CTGGTGGAGGCCAAAAACATGGTGATGAGTTTTCGAGTCTCCGACCTTCAGATGCTCCTGGGTTTC
GTGGGCCGGAGTAAGAGTGGACTGAAGCACGAGCTCGTCACCAGGGCCCTCCAGCTGGTGCAGTTT
GACTGTAGCCCTGAGCTGTTCAAGAAGATCAAGGAGCTGTACGAGACCCGCTACGCCAAGAAGAAC
TCGGAGCCTGCCCCACAGCCGCACCGGCCCCTGGACCCCCTGACCATGCACTCCACCTACGACCGG
GCCGGCGCTGTGCCCAGGACTCCGCTGGCAGGCCCCAATATTGACTACCCCGTGCTCTACGGAAAG
TACTTAAACGGACTGGGACGGTTGCCCGCCAAGACCCTCAAGCCAGAAGTCCGCCTGGTGAAGCTG
CCGTTCTTTAATATGCTGGATGAGCTGCTGAAGCCCACCGAATTAGTCCCACAGAACAACGAGAAG
CTTCAGGAGAGCCCGTGCATCTTCGCATTGACGCCAAGACAGGTGGAGTTGATCCGGAACTCCAGG
GAACTGCAGCCCGGAGTTAAAGCCGTGCAGGTCGTCCTGAGAATCTGTTACTCAGACACCAGCTGC
CCTCAGGAGGACCAGTACCCGCCCAACATCGCTGTGAAGGTCAACCACAGCTACTGCTCCGTCCCG
GGCTACTACCCCTCCAATAAGCCCGGGGTGGAGCCCAAGAGGCCGTGCCGCCCCATCAACCTCACT
CACCTCATGTACCTGTCCTCGGCCACCAACCGCATCACTGTCACCTGGGGGAACTACGGCAAGAGC
TACTCGGTGGCCCTGTACCTGGTGCGGCAGCTGACCTCATCGGAGCTGCTGCAGAGGCTGAAGACC
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AGCGAGATCGCCACCACCGGTGTGCGGGTGTCCCTCATCTGTCCGCTGGTGAAGATGCGGCTCTCC
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AACGAGAAGAAGCCCACCTGGATGTGCCCCGTGTGCGACAAGCCAGCCCCCTACGACCAGCTCATC
ATCGACGGGCTCCTCTCGAAGATCCTGAGCGAGTGTGAGGACGCCGACGAGATCGAGTACCTGGTG
GACGGCTCGTGGTGCCCGATCCGCGCCGAAAAGGAGCGCAGCTGCAGCCCGCAGGGCGCCATCCTC
GTGCTGGGCCCCTCGGACGCCAATGGGCTCCTGCCCGCCCCCAGCGTCAACGGGAGCGGTGCCCTG
GGCAGCACGGGTGGCGGCGGCCCGGTGGGCAGCATGGAGAATGGGAAGCCGGGCGCCGATGTGGTG
GACCTCACGCTGGACAGCTCATCGTCCTCGGAGGATGAGGAGGAGGAGGAAGAGGAGGAGGAAGAC
GAGGACGAAGAGGGGCCCCGGCCCAAGCGCCGCTGCCCCTTCCAGAAGGGCCTGGTGCCGGCCTGC

Figure 6 (continued)

>PIASy(prey)
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GTGGGCCGGAGTAAGAGTGGACTGAAGCACGAGCTCGTCACCAGGGCCCTCCAGCTGGTGCAGTTT
GACTGTAGCCCTGAGCTGTTCAAGAAGATCAAGGAGCTGTACGAGACCCGCTACGCCAAGAAGAAC
TCGGAGCCTGCCCCACAGCCGCACCGGCCCCTGGACCCCCTGACCATGCACTCCACCTACGACCGG
GCCCGGCGCTGTGCCCAGGACTCCGCTGGCAGGCCCCAATATTGACTACCCCGTGCTCTACGGAAAG
TACTTAAACGGACTGGCACGGTTGCCCGCCAAGACCCTCAAGCCAGAAGTCCGCCTGGTGAAGCTG
CCGTTCTTTAATATGCTGGATGAGCTGCTGAAGCCCACCGAATTAGTCCCACAGAACAACGAGAAG
CTTCAGGAGAGCCCGTGCATCTTCGCATTGACGCCAAGACAGGTGGAGTTGATCCGGAACTCCAGG
GAACTGCAGCCCGGAGTTAAAGCCGTGCAGGTCGTCCTGAGAATCTGTTACTCAGACACCAGCTGC
CCTCAGGAGGACCAGTACCCGCCCAACATCGCTGTGAAGGTCAACCACAGCTACTGCTCCGTCCCG
GGCTACTACCCCTCCAATAAGCCCGGGGTGGAGCCCAAGAGGCCGTGCCGCCCCATCAACCTCACT
CACCTCATGTACCTGTCCTCGGCCACCAACCGCATCACTGTCACCTGGGGGAACTACGGCAAGAGC
TACTCGGTGGCCCTGTACCTGGTGCGGCAGCTGACCTCATCGGAGCTGCTGCAGAGGCTGAAGACC
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AGCGAGATCGCCACCACCGGTGTGCGGGTGTCCCTCATCTGTCCGCTGGTGAAGATGCGGCTCTCC
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AACGAGAAGAAGCCCACCTGGATGTGCCCCGTGTGCGACAAGCCAGCCCCCTACGACCAGCTCATC
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GGCAGCACGGGTGGCGGCGGCCCGGTGGGCAGCATGGAGAATGGGAAGCCGGGCGCCGATGTGGTG
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>PLIP
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GCCTCGGTTTTTCCCCAGAATGGAGCCGCCCGTAGGGCAGTGGCAGCCCAGCCAGGACGGAAGCGA
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CGCATGACTGGCAGCCTGGTGTCTGATCGAAGCCACGACGACATCGTCACCCGGATGAAGAACATT
GAGTGCATTGAGCTGGGCCGGCACCGCCTCAAGCCGTGGTACTTCTCCCCGTACCCACAGGAACTC
ACCACATTGCCTGTCCTCTACCTGTGCGAGTTCTGCCTCAAGTACGGCCGTAGTCTCAAGTGTCTT
CAGCGTCATTTGACCAAGTGTGACCTACGACATCCTCCAGGCAATGAGATTTACCGCAAGGGCACC
ATCTCCTTCTTTGAGATTGATGGACGTAAGAACAAGAGTTATTCCCAGAACCTGTGTCTTTTGGCC
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ACACTGTCAGAGGACATCGTGGATGGCCATGAGCGGGCCATGCTCAAGCGGCTCCTGCGGATCGAC
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Figure 6 (continued)

>PTN
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CTCGCCCCTTGCAACAAAGGCAGACTGAGCGCCAGAGAGGACGTTTCCAACTCAAAAATGCAGGCT
CAACAGTACCAGCAGCAGCGTCGAAAATTTGCAGCTGCCTTCTTGGCATTCATTTTCATACTGGCA
GCTGTGGATACTGCTGAAGCAGGGAAGAAAGAGAAACCAGAAAAAAAAGTGAAGAAGTCTGACTGT
GGAGAATGGCAGTGGAGTGTGTGTGCCCACCAGTGGAGACTGTGGGCTGGGCACACGGGAGGGC
ACTCGGACTGGAGCTGAGTGCAAGCAAACCATGAAGACCCAGAGATGTAAGATCCCCTGCAACTGG
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>PTPK
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>SETBD1
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CCTGTGAAGCCTGAAGGACTTCGCCGCCCACCTAGTAAGACTAGTATGCATCAAAGCCGAAGACTC
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ATGAAGCGTCAAGTGGCAGTAAAATCAACCCGAGGCTTTGCTCTTAAATCAACCCATGGGATTGCA
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TACCTCAACCACAGTTGCAGCCCCAACCTGTTTGTCCAGAATGTCTTCGTGGATACCCATGATCTT
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AGAGGACGTCTTCTT

Figure 6 (continued)

>SH3GL3
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CTGGCAGGGCACAGTTTC
>SUMO-3
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GAACGACAGGGATTGTCAATGAGGCAGATCAGATTCCGATTTGACGGGCAACCAATCAATGAAACA
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GGTGTCTAC
>TAL1
AGCTCACCCGTGAAGCGTCAGAGGATGGAGTCCGCGCTGGACCAGCTCAAGCAGTTCACCACCGTG
GTGGCCGACACGGGCGACTTCCACGCCATCGACGAGTACAAGCCCCAGGATGCTACCACCAACCCG
TCCCTGATCCTGGCCGCAGCACAGATGCCCGCTTACCAGGAGCTGGTGGAGGAGGCGATTGCCTAT
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ACCGACAAGAAATCCTATGAGCCCCTGGAAGACCCTGGGGTAAAGAGTGTCACTAAAATCTACAAC
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AACGCCAAGCTGGTGCCTGTGCTCTCAGCCAAGGCGGCCCAAGCCAGTGACCTGGAAAAAATCCAC
CTGGATGAGAAGTCTTTCCGTTGGTTGCACAACGAGGACCAGATGGCTGTGGAGAAGCTCTCTGAC
GGGATCCGCAAGTTTGCCGCTGATGCAGTGAAGCTGGAGCGGATGCTGACAGAACGAATGTTCAAT
GCAGAGAATGGAAAG

Figure 6 (continued)

>TCPG
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GATGATGTTGGAACAGGAGCAGGCCTGTTGGAAATCAAGAAAATTGGAGATGAATACTTTACTTTC
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GCTCCTGATGCTGGCCAGGAG
>VIM
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TCCGGGAGCCAGTCCGCGCCACCGCCGCCGCCCAGGCCATCGCCACCCTCCGCAGCCATGTCCACC
AGGTCCGTGTCCTCGTCCTCCTACCGCAGGATGTTCGGCGGCCCGGGCACCGCGAGCCGGCCGAGC
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CTGCGGAGCAGCGTGCCCGGGGTGCGGCTCCTGCAGGACTCGGTGGACTTCTCGCTGGCCGACGCC
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GAA

Figure 6 (continued)

>VIMc
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>ZHX1
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TCAGATGAC

Figure 6 (continued)

>ZNF33B
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Figure 6 (continued)

Nucleotide sequence data (fasta format)

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>GAPD
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Protein sequence data (fasta format)

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Figure 6 (continued)

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Figure 6 (continued)

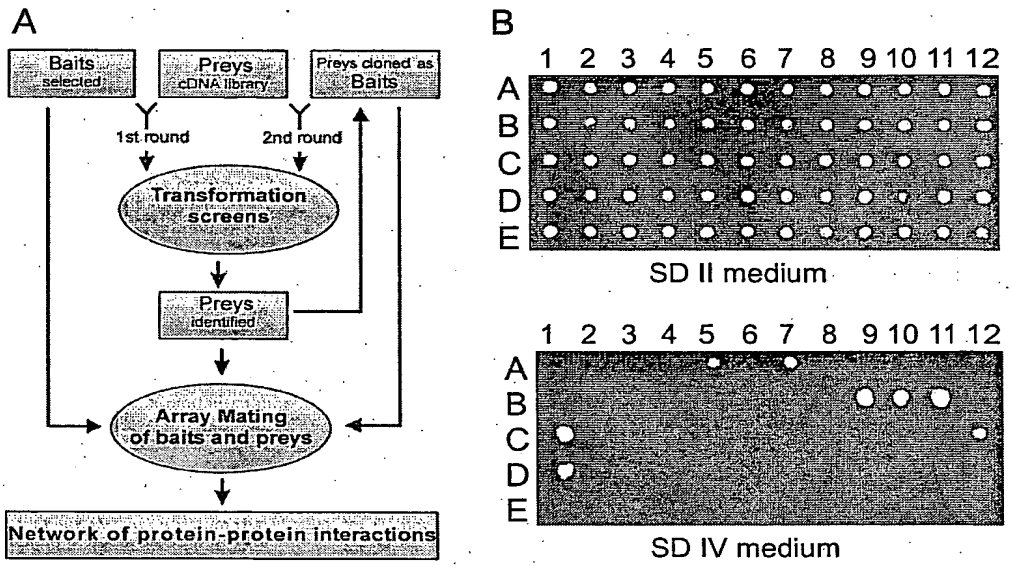
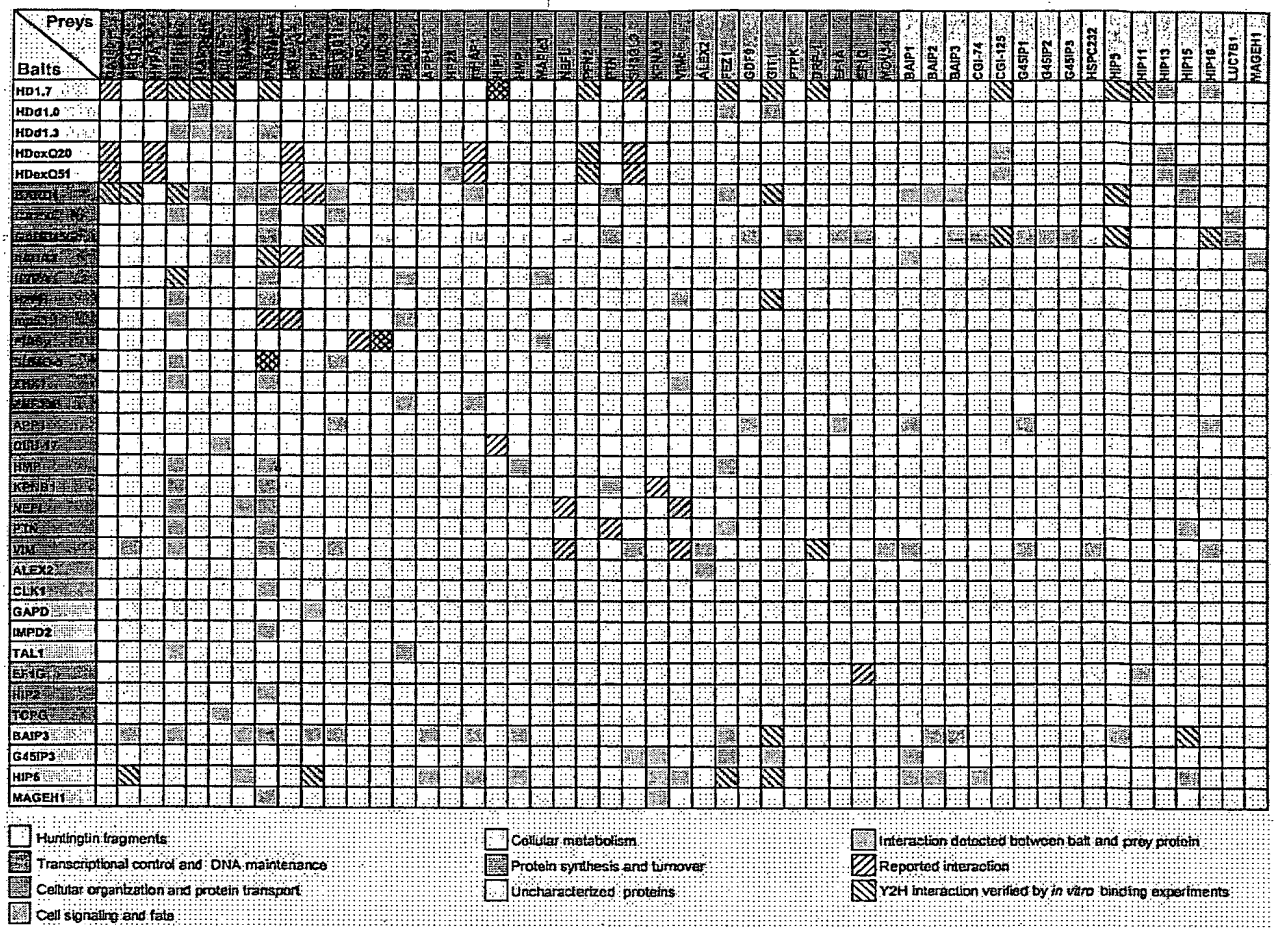


Figure 7

A



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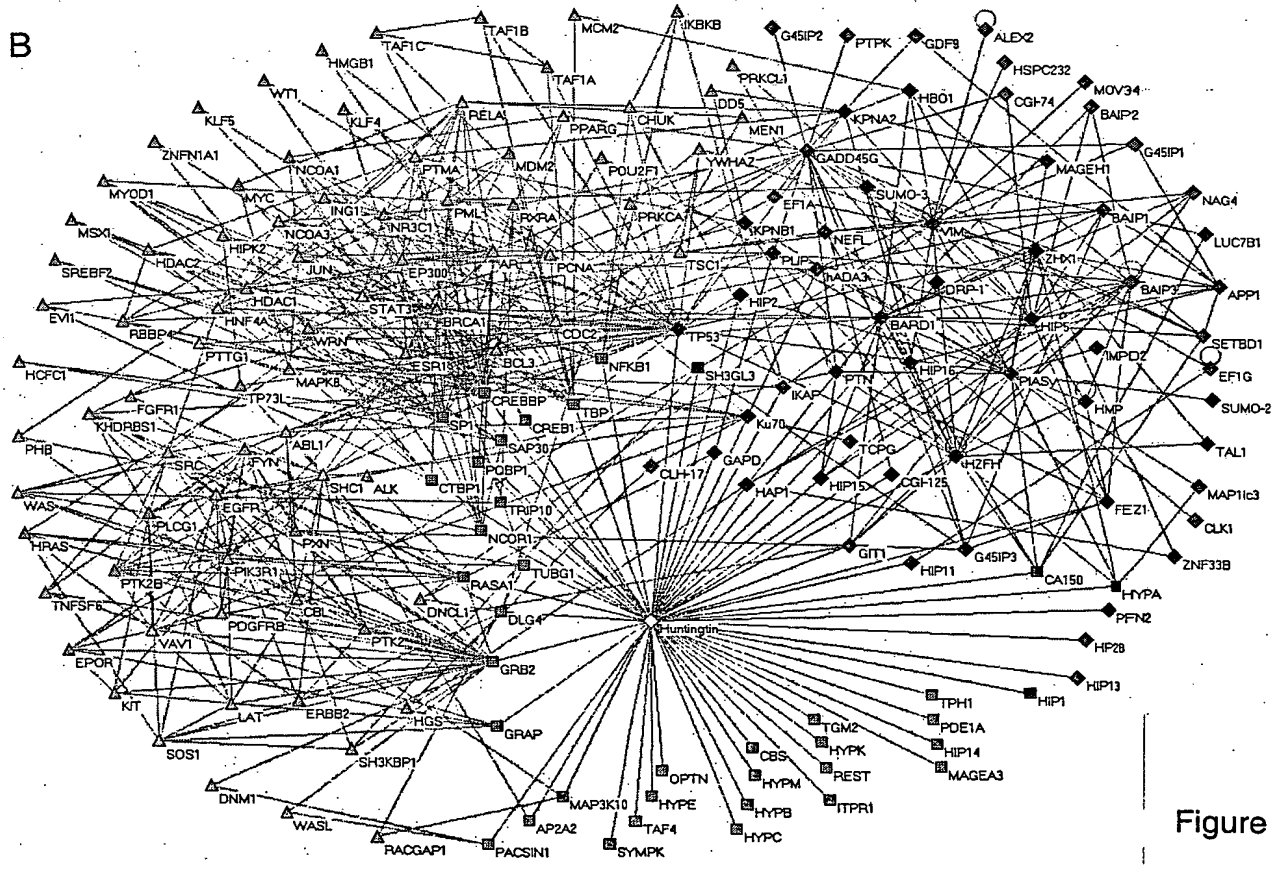

Figure 8

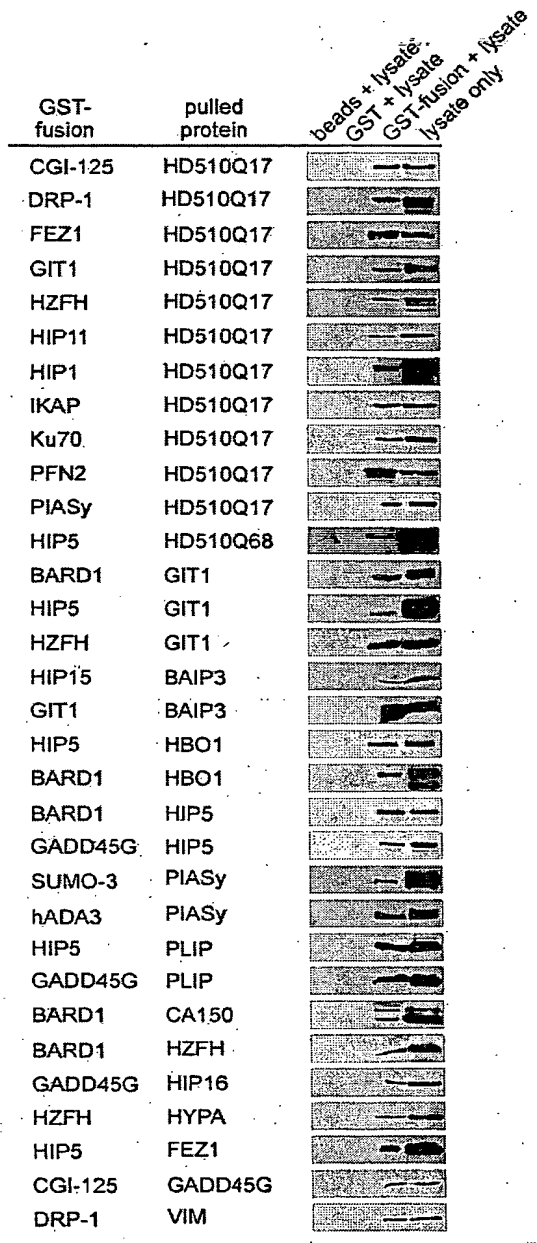

Figure 9

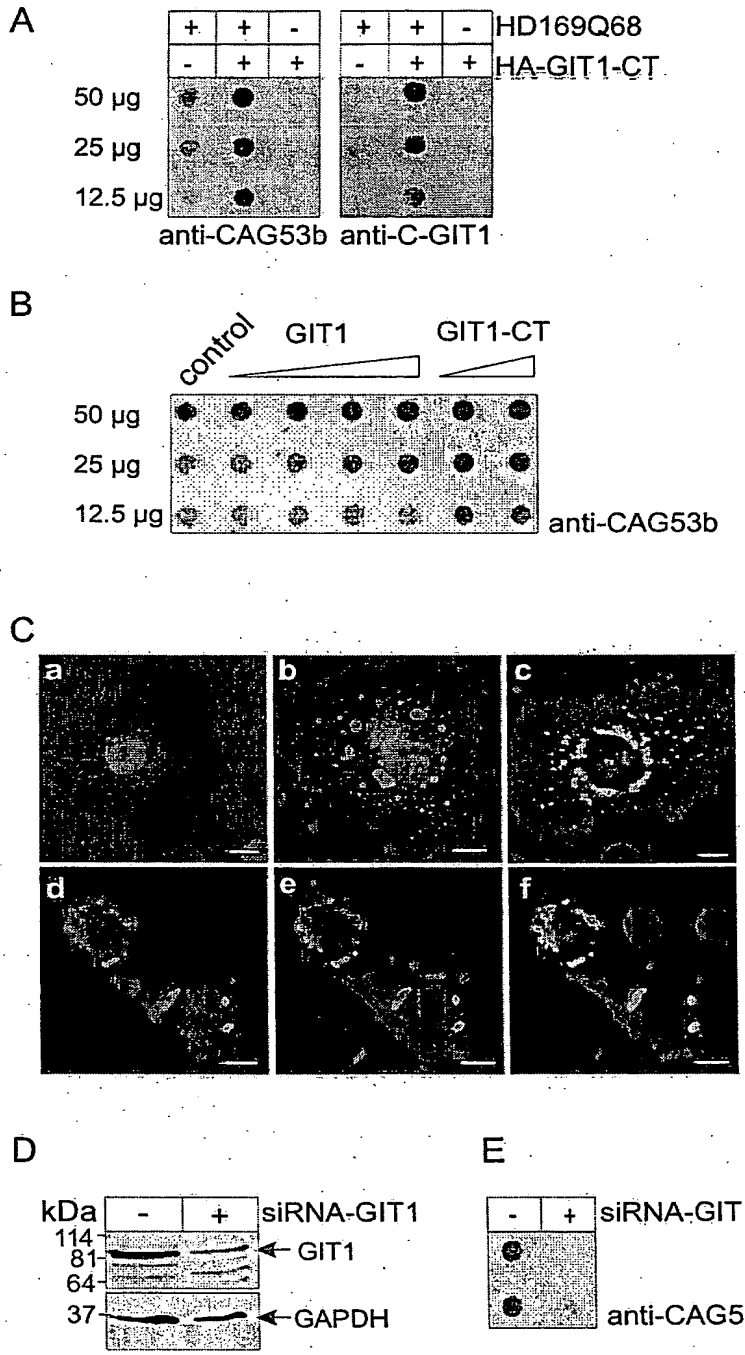

Figure 10

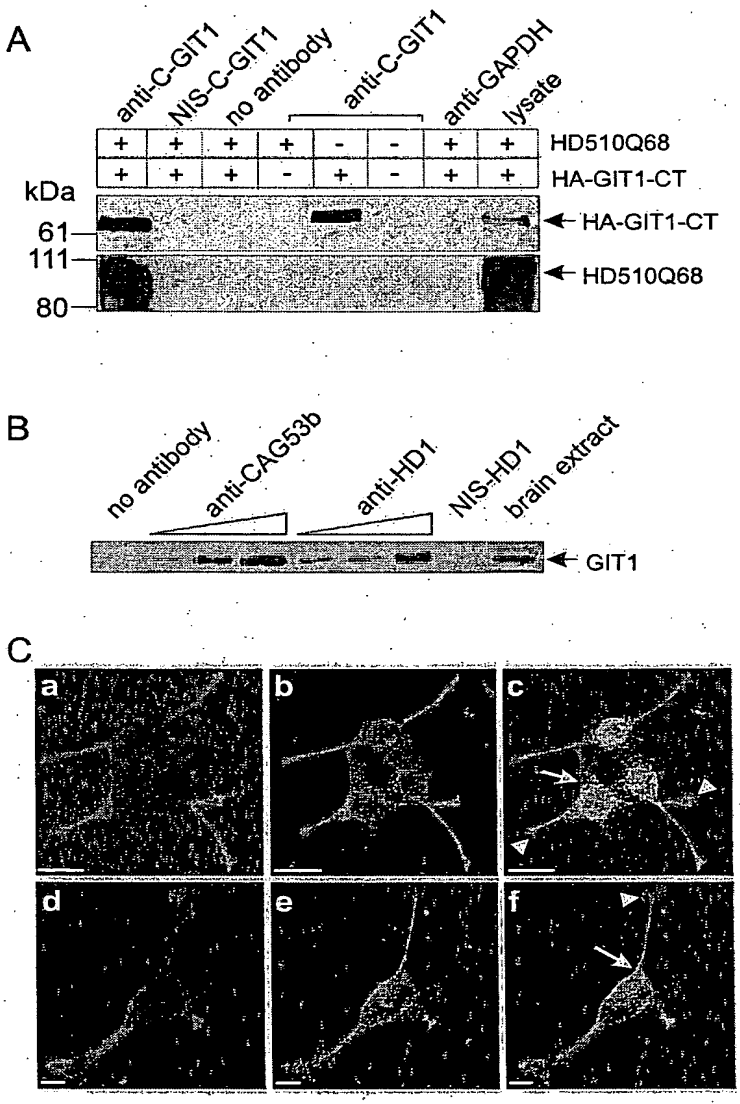

Figure 11

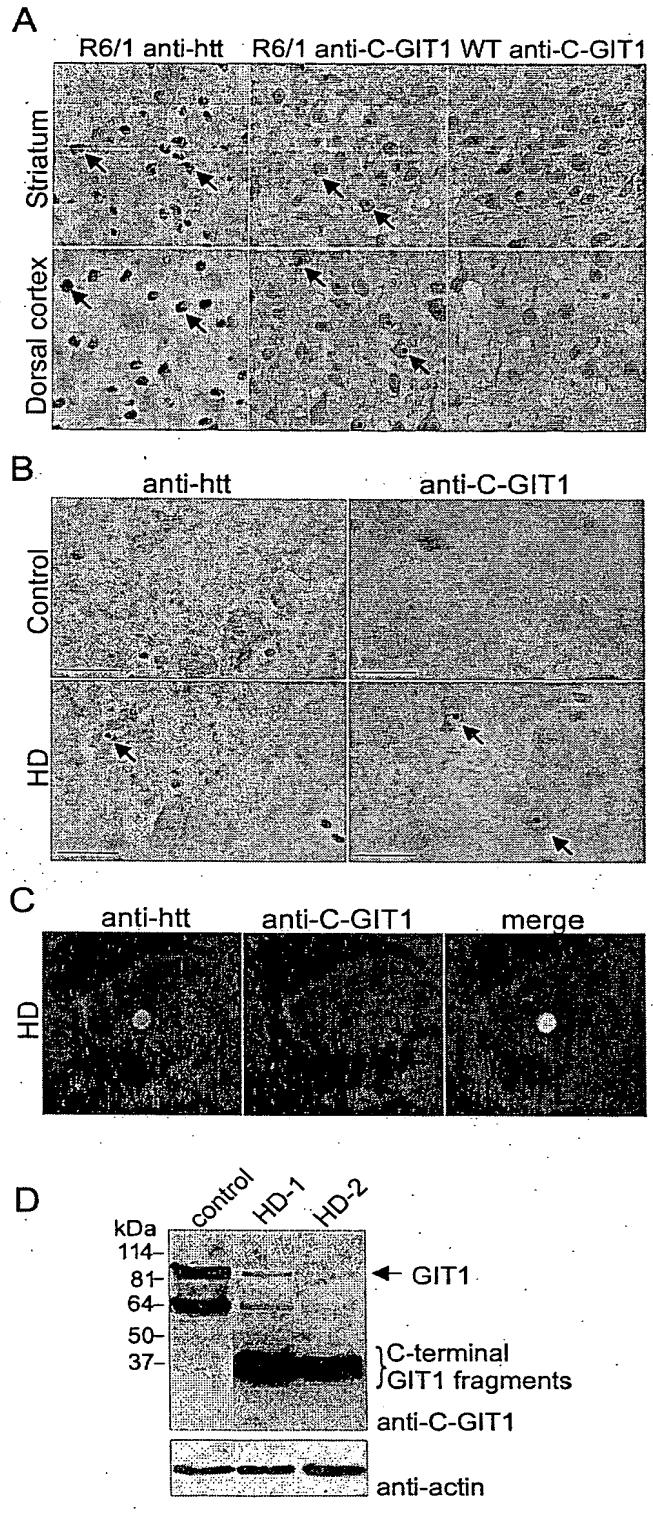

Figure 12

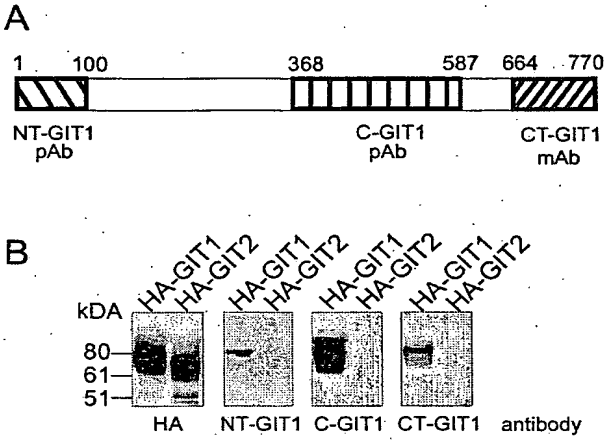

Figure 13